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ALIGNMENTS

JOURNAL COMMENT KEYWORDS SOURCE ORGANISM RESULT 1 BQ636767 LOCUS REFERENCE DEFINITION ACCESSION VERSION TITLE AUTHORS BQ636767 he01b10.y1 Human Retina c heono sapiens cDNA clone h BQ636767 BQ636767.1 GI:21761226 Wistow,G., Bernstein,S.L., Wyatt,M.K., Ray,S., Behal,A., Touchman, J.W., Bouffard,G., Smith,D. and Peterson,K. Expressed sequence tag analysis of human retina for the NEIBank Project: Retbindin, an abundant, novel retinal cDNA and alternative splicing of other retinal-preferred gene transcripts Mol. Vis. 8 (4), 20021 In press Tel: 301 402 3452 Fax: 301 496 0078 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 334) National Eye Institute 6/331, NIH, Bethesda, MD 20892-2740, Section on Molecular Structure and Function Homo sapiens Homo sapiens (human) graeme@helix.nih.gov 01 row: b column: 10 334 bp mRNA linear EST 15-JUL-2002 cDNA (Un-normalized, unamplified): hd/he he01b10 5', mRNA sequence. the NEIBank and alternative

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JOURNAL COMMENT
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            Email: yongsung@mail.kribb.re.kr
Plate: 51 row: E column: 10
High quality sequence stop: 355.
                                                                                                                                                            Genome Research Center
                                                                                                                                                                                                                Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R., Oh,K.J., Cheong,J.E., Sohn,H.Y., Kim,J.M., Park,H.S., Kim,S. and Kim,Y.S.
                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 355)
Kim,N.S., Hahn,Y., Oh,J.H., Lee,J.Y., Ahn,H.Y., Chu,M.Y., Kim,M.R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mRNA sequence
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                                                                                                                                  Korea Research Institute of Bioscience & Biotechnology
                                                                                                                                                                                    Contact: Kim YS
                                                                                                                                                                                                     Unpublished
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                             Eoeun-dong Yuseong-gu,
l: +82-42-860-4470
x: +82-42-860-4409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: Eye; Vector: pSPORT1; Neural retina tissue was dissected from two 80 year old donors with no observed eye dissease. 100ug of total RNA was used for library construction. A directionally cloned cDNA library in the pSPORF1 vector (Life Technologies) was constructed at Bioserve Biotechnology (Laurel MD) essentially following the protocols of the SuperScript Plasmid System full details of which are contained in the manufacturer's Instruction manual (http://www.lifetech.com/). First strand synthesis was carried out using a Not I primer-adapter [5'.pGACTAGTTCTAGATCGGGACGGCCCCT]15-3']. EST analysis was performed on the unamplified library at the NIH Intramural Sequencing Center (NISC)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="Retina"
/dev_stage="Adult"
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/mol_type="mRNA"
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Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
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                                                       3201 Carnegie Ave, Cleveland, Tel: 216 431 9900 Fax: 216 361 9596
                                                                                                                                                                           Contact: Scott J. Cain
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/organism="Homo_sapiens"
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Pred. No. 5.2e-45;
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                                                                                                                                                                                                      Email: johng@tigr.org
Plate: 149
                                                                                                                                                                                                                                   9712 Medical Center Dr.,
Tel: 301 838 3528
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                          Unpublished
Contact: John Quackenbush
                                                                                                                                                                                                                                                                                                                     Assessment of gene expression patterns in a model of metastasis using a 19,200 element cDNA microarray
                                                                                                                                                                                                                                                                                                                                                                            Hegde, P., Qi, R., Abernathy, K.,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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ilarity 100.0%;
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/note-"See 'Creation of Genome-wide Protein Expression
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
                                                                    /db_xref="taxon:9606"
/Clone_lib="MAGE resequences, MAGF"
/note="vector: pBluescriptsKm"
153 c 137 g 107 t
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Mismatches 0;
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Pred. No. 6e-45;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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601070792F1 NIH_MGC_12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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BE543478.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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1 300 c 247 g 88 t
                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nlh.gov
Tissue Procurement: Elias Campo, M.D., Michael R.
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National Cancer Institute, Cancer Genome Anat
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DNA Sequencing by: Washington University Genome Sequencing (
Clone distribution: NCI-CGAP clone distribution information found through the I.M.A.G.E. Consortium/LLNL at:
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/clone_lib="NCI_CGAP_CO3"
/note="Vector: pm7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; lst strand cDNA was prepared from 12 pooled bulk tumor samples and primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT773 vector. Library went through one round of normalization. "
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/clone="IMAGE:926191"
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/mol_type="mRNA"
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nf80b04.x5 |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arraying: Grey Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:926191"
/sex="pooled"
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/note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNP was prepared from 12 pooled bulk tumor samples and primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. "
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result Score Query Match 100.0 45 163396 176584 183099 188540 192093 230493 242175 Length 482 5635 368 368 368 368 368 379 179 179 168 182 182 182 182 186 388 398 398 402 364 DВ 966667 909 10 6 6 6 6 6 12 6 SYNECOPST BC025790 AR270687 AX332609 AX333248 AX411203 BD160949 SYNPSTIAA A24907 E03479 AC011108 AC121205 AL837521 MMMPGC60 AL161445 AC027812 AC034144 AR166138 E13091 AR166137 E13090 AF048700 AR028526 AR075817 BD007521 MMPIMP12 E01725 AC096463 I08731 A24905 A11350 A11352 E01574 E02456 SSPEC60 E08411 E03120 HSPSTI S46866 AR059723 AX014851 ASVPSKAN8 Y00705 Homo sapien X06342 Mouse mRNA E01725 Synthetic D E02456 DNA encodin M31103 Synthetic h ALI61445 Human DNA AC027812 Homo sapi AC034144 Homo sapi AC011108 Homo sapi AC011108 Idomo sapi AC121205 Rattus no AL837521 Mouse DNA AC096463 Rattus no 108731 Sequence 19 A11350 Artificial A11352 Artificial A11352 Artificial A24905 PSTI master M21832 Synthetic h A24907 DSNA encodi X84307 Artificial AF048700 Homo sapi AR028526 Sequence AR075817 Sequence BD007521 Tumor-ass BC025790 Homo sapi AR270687 Sequence S46866 PEC-60=gast X67109 S.scrofa PE Y11505 M.musculus AR166138 Sequence E13091 Human cDNA AX014851 Sequence M11949 Human E01574 cDNA sequen M22196 Synthetic E AR059723 Sequence E08411 DNA sequenc AR166137 Sequence E13090 Human an cD E01819 DNA sequenc E03120 DNA sequenc Description ното зарі

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ALIGNMENTS

RESULT 1
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ACCESSION
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KEYWORDS SOURCE ORGANISM FEATURES REFERENCE AUTHORS TITLE 1 (bases 1 to 180) Kato, S. Yamaguchi, T., Sekine, S. and Kamata, Human PEC-60-like protein and DNA encoding t Patent: US 6280968-A 1 28-AUG-2001; AR166137 180 bp Sequence 1 from patent US 6280968. AR166137 AR166137.1 Unclassified. Unknown Unknown Location/Qualifiers GI:16241312 DNA the linear same PAT 17-OCT-2001

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ORIGIN
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent: JP 1997124698-A
SAGAMI CHEM RES CENTER
OS HOMO Sapiens (human
PN JP 1997124698-A/1
PD 13-MAY-1997
PF 27-OCT-1995 JP 1995
PI KATO MASASHI, YAMAG
PC C12P21/02, C07H21/0
PC C12P21/02, C12R1:19
PC (C12P21/02, C12R1:19
PC (C12P21/02, Linear;
FH Key Looplogy: Linear;
FH Key CHEM
PC TYPEN CONTEN CONTE
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Kato, M., Yamaguchi, T., Sekine, S. and Kamata, M. HUMAN PEC-60LIKE PROTEIN AND DNA CAPABLE OF CODING THE SAME Patent: JP 1997124698 A 1 13-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP 1997124698-A/1.
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E13090
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                                                                    GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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27-OCT-1995 JP 1995280272
KATO MASASHI, YAMAGUCHI TOMOKO, SEKINE SHINGO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
JP 1997124698-A/1
                                                                                                                                                                                                                           Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
49 c 40 g 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .180
/organism="unknown"
49 c 40 g
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/tissue_type='stomach cancer'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product='PEC-60-like
                                                                                                                                                                                                                Score 180; DB 6;
Pred. No. 2.6e-47;
; Mismatches 0;
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Pred. No. 2.6e-47;
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PEC-60-like protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (17-FEB-1998)
Yue-Yang Road, Shanghai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human gastrointestinal
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                                                                                                                                                                                                                                                  h 100.0%;
Similarity 100.0%;
80; Conservative (
                                                                                                                                                                                                         CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
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rcsomsnivcgtdgltytnecqlclariktkqdiqimkdgkc"
97 c 98 g 72 t
                                                                                                                                                                                                                                                                                                                                         /codon_start=1
/product="gastrointestinal peptide"
/protein_id="AACOS124.1"
/db_xref="G1:2935440"
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                               /gene="PEC-60"
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                                                    from patent US
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Pred. No. 2.6e-47;
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RESULT 6
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Matches 180; Conserv
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BD007521
BD007521.1 GI:18635894
JP 2001503629-A/1
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Bandman,O., Goll,S.K. and Murry,L.E.
Tumor-associated Kazal inhibitor
Patent: US. 4958699-A 2 28-SEP-1999;
Location/Qualifiers
1. .388
                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
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Patent: US 5858
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Bandman, O., Goli, S.K. and Murry, L.E.
Tumor-associated kazal inhibitor-like
                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
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Location/Qualifiers
1. 388
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105 c 102 g
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105 c . 102 g
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Pred. No. 2.6e-47;
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AUTHORS
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PN JP 2001503629-A/1
PD JP 605.NOV-1997 JP 1998521770
PR 06-NOV-1996 US 08/744670
PR 06-NOV-1996 US 08/744670
PR 06-NOV-1996 US 08/744670
PC C12N15/09, A61K38/55,A61K39/395,A61P1/00,A61P35/00,C07K14/81,
PC C07K16/38,
PC C12N1/15,C12N1/21,C12N5/10,C12Q1/68,G01N33/574, PC C12N15/00,C12N5/00,
PC A61K37/64
CC A61K37/64
FT Source /organism='Unidentified'.
FT Source /organism='Unidentified'.
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Sequence 2 :
AR166138
AR166138.1
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Bandman,O., Goli,S.K. and Murry,L.E.
Bandman,O., Goli,S.K. and Murry,L.E.
Tumor-associated KAZAL type inhibitor
Patent: JP 2001503629-A 1 21-MAR-2001;
INCYTE PHARMACEUTICALS INC
                                                                                                                Kato.S., Yamaguchi,T., Sekine,S. and Kamata,
Human PBC-60-like protein and DNA encoding (
Patent: US 6289958-A 2 28-AUG-2001;
Location/Qualifiers
                                                                                                                                                                                                       Unknown.
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               Similarity
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JP 2001503629-A/1
21-MAR-2001
05-NOV-1997 JP 1998521770
06-NOV-1996 US 08/744670
OCAN BANDMAN, SURYA K GOLI, LYNN E MURRY
C12N15/09, A61K38/55, A61K39/395, A61P1/00, A61P35/00, C07K14/81,
C07K16/38
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110 c 105 g
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
105 c 102 g 7
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1. .388
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Score 180; DB 6;
Pred. No. 2.6e-47;
; Mismatches 0;
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Pred. No. 2.6e-47;
Mismatches 0;
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1 GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT 60

PAT 07-SEP-2000

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RESULT 8
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PN JP 19
PN JP 19
PN 13-MA
PF 27-OC
PF KATO
MITSUHISA
PC C07K1
PC C12P2
PC (C12P2
PC (C1
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HUMAN PEC-60LIKE PROTEIN AND DNA CAPABLE OF CODING THE SAME
Patent: JP 1997124607EIN AND 13-MAY-1997;
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                                                                                                              CTCTGCTTGGCCCGGATAAAAACCCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                               GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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O,M., Yamarran
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                                                                                                                                                                                                                                                                                                                                 107
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KATO MASASHI, YAMAGUCHI TOMOKO, SEKINE SHINGO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-OCT-1995 JP 1995280272
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                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
110 c 105 g 7
                                                                                                                                                                                                                                                                                                                                                                                                                       /product='PEC-60-like Location/Qualifiers
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/tissue_type='stomach ca
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Pred. No. 2.6e-47;
; Mismatches 0;
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Best Local S
Matches 180
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JOURNAL
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                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Cetartiod
1 (bases 1 to 364)
Metsis,M., Cintra,A., Solfrin.
Morrasutti,D.G., Ostenson,C.G
et,al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                          GenBank staff at the National Library of Medicine created this entry \{NCBI\ gibbsq\ 115614\} from the original journal article. This sequence comes from Fig. 1.
                                                                                                                                                               Molecular cloning of PEC-60 and expression of its in the gastrointestinal tract and immune system J. Biol. Chem. 267 (28), 19829-19832 (1992) 93015834
                                                                                                                                                                                                                                                                                                                                                                                                  $46866
PEC-60-gastrointestinal
$46866
$46866.1 GI:257597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human nucleic acid sequences from ovarian tumour tissue Patent: WO 9953040-A 41 21-OCT-1999; SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (DE); HINZMANN BERND (DE); ROSEWITHAL ANDRE (DE); METAGEN GES FUER GENOMFORSCHUN (DE); PILARSKY CHRISTIAN (DE)
LOCATION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                 Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                Sus scrofa
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
111 c 107 g 7
/organism="Sus scrofa"
/mol_type="mRNA"
/db_xref="taxon:9823"
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                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                      364 bp
peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA
[swine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hinzmann, B.,
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MAM πRNA,

05-AUG-1999 364 nt].

303 180 243

mRNA and

peptide

Euteleostomi;
Sus.

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SOURCE
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ORIGIN
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                                                                                                                                                                                                                                                                                         AUTHORS
TITLE
  Query Match
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                                                                                                                                                                                                                                                                                                                                                       pec-60 gene.
Sus scrofa (pig)
                                                                                                                                                                                                                                                                 Submitted (22-JUN-1992) M. of Molecular Neurobiology,
                                                                                                                                                                                                                                                                                                                                                                                             S.scrofa PEC-60 mRNA
X67109
                                                                                                                                                                                                                                                          Stockholm,
                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
                                                                                                                                                                                                                                                                                                                                                Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                 X67109.1 GI:2033
                                                                                                                                                                                                                                                                                                                                                                                                                  SSPEC60
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                                                                                                                                                                                                                                                                                                      Metsis,M.
                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria;
1 (bases 1 to 364)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="gastrointestinal peptide; This sequence Fig. 1; Author includes translated amino acids putative initiation codon"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /translation="MAVRLWVVALALAALFIVDREVPVSAEKQVFSRMPICEHMTESP
DCSRIYDPVCGTDGVTYESECKLCLARIENKQDIQIVKDGEC"
93 c 108 g 72 t
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                                                       /note='
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                                                                                                                       /product="peptide PEC-60"
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/db_xref="GI:2034"
                                                                                                                                                                                                 /organism="Sus scrofa"
/mol_type="mRNA"
/db_xref="taxon:9823"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23. .283
                                                                                                                                                       /evidence=experimental
                                                                                                                                                                 /codon_start=1
                                                                                                                                                                                        /clone_lib="pig duodenum cDNA library
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/db_xref="GI:5705936"
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81.7%;
  70.78;
                                                                  peptide PEC-60"
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Pred. No. 3.7
  Score 127.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                   364
                                                                                                                                                                                                                                                                   Metsis, Dept. of Medical Chemistry, Lab
Karolinska Institute, Box 604 00, 10401
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Best Local Similarity 81.7 Matches 147; Conservative
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143; Conser
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Y11505
Y11505.1 GI:2344863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Krause,R., Hemberger,M., Messerschmid,M., Kothary,R. and Fundele,R. Molecular cDNA cloning and expression of murine Mpgc60, a gene predominantly expressed in the intestinal tract
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (26-FEB-1997) R. Krause, Ropers, Ihnestr. 73, 14195 Berlin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Krause, R.
Direct Submission
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                           /product="Mpgc60 protein"
/protein_id="CAA72282.1"
/protein_id="CAA72282.1"
/db_xref="GI:2344864"
/db_xref="SPTREMBL:035679"
/translation="MAMHLWLYTLTLGPLLGMDRELMVSAGSLVFPRMPFCEHMAELP
/translation="MAMHLWLYTLTLGPLLGMDRELMVSAGSLVFPRMPFCEHMAELP
NCPQTPNLICGTDGLTYENECHLCLTRMKTMKDIQIMKDGQC"
129 c 136 g 95 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                    /sub_strain="Hannover (random bred)"
/db_xref="taxon:10090"
/tissue_type="intestine"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/strain="NMRI"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://www.sanger.ac.uk/HGP/Chr9
This sequence is the entire insert of clone RP11-326F20 The true left end of clone RP11-34B24 is at 100362 in this sequence. The true right end of clone RP11-54K16 is at 54767 in this sequence. This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. RP11-326F20 is from the literature of the result of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (04-DEC-2001) Wellcome Trust Sanger Institute, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail enquiries:
humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
on Nov 10, 2000 this sequence version replaced gi:11071610.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AL161445

163396 bp DNA linear PRI 04-DEC-2001

Human DNA sequence from clone RPII-326F20 on chromosome 9 Contains

the GGTA1 gene for glycoprotein alpha-galactosyl transferase 1, the

SPINK4 gene for Kazal type 4 serine protease inhibitor, the 3' end

of the BAG1 gene for BCL2-associated athanogene and a CpG island,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the library RPCI-11.2 constructed by the group of Pieter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Group.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
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GGTA1; glycoprotein; Kazal; s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 9, constructed by the Sanger Centre Chromosome 9 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Em:, EMBL; Sw:, SWISSPROT; Tr:, TREMBL; Wp:, WORMPEP; on the WORMPEP database can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      variation annotation may not be found in the sequence submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pBACe3.6.
                                                                                         /note="MER44A repeat: matches 225.
440. .555
                                                                                                                                                                                                                                                               /note="AluY repeat: matches 77. .309 of consensus" 239. .314
                                                                                                                                                                              /note="L1M4 repeat: matches 2591.
331. .438
                                                                                                                                                                                                                                                                                                                                                             /clone_lib="RPCI-11.2"
5. .237
                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="RP11-326F20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:9606"
/chromosome="9"
                                        'note="AluSq/x repeat: matches 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         location/Qualifiers
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serine protease inhibitor; SPINK4.
                                                                                                                                                                                                                  . 2666
                                        .126 of consensus'
                                                                                                                             .331 of consensus"
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3046. .3
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8784 . . .
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                                                                                     /note="MIR repeat: matches 112. .257 of consensus"
complement(14283. .70891)
                                                                                                                                                                                                                                                                     /note="match: GSS: Em:AQ210501" complement(12571 .12678)
                                                                                                                                                                                                                                                                                                        /note="9 copies 10 mer gtgtgtgtt 72% conserved"
complement(12461. .12790)
                                                                                                                                                                                                                                                                                                                                                                                 /note="MIR repeat: matches 35. 12243. .12346
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="MIR repeat: matches 132. .252 of consensus" 11874. .12110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="AluSp repeat: matches 1. .302 of consensus" 9086. .9106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="AluSx repeat: matches 1..304 of consensus"
8770..8783
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7523. .7596
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1982. .2181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="AluJo/FRAM repeat: matches 158. .298 of consensus' 795. .1193
                                                   complement(join(14283. .17232,17420.
                                                                                                                            14033.
                                                                                                                                                               /note="FLAM_A repeat: matches 1.
13290. .13517
                                                                                                                                                                                                  /note="MER21B repeat: matches 232.
13171. .13289
                                                                                                                                                                                                                                                                                                                                               /note="52 copies 2 mer tg
12254. .12343
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12121. .12219
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1194. .1491
                                                                                                                                                                                                                                                    note="match: GSS: Em:AQ177335"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9152. .9751
/note="L1MB3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="L1P1 repeat: matches 5753. .5825 of consensus"
/product="bA326F20.1 (glycoprotein, alpha-galactosyl
                                                                                                                                         'note="MER21B repeat: matches 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'note="MIR repeat: matches 61.
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                                 .24252,38835.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    matches 1. .304 of consensus"
                                   .39070,70404. .70891))
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                                                   .17524,19637. .19759,
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                                           67
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                    CTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGTTCCCAG 66
                                                                                                                                                                             Conservative
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/note="L2 repeat: matches 2241. .2411 of consensus"
19126. .19243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /evidence=not_experimental
/product="ba326F20.1 (g1ycoprotein, alpha-galactosyl
transferase 1)"
/protein_id="CAD13306.1"
/db_xref="G1:17384434"
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RDLSRLPQLVGVSTPLQGGSNSAAAIGQSSGELRTGGARPPPPLGASSQPRPGGDSSP
VVDSGPGPASNLTSVPVPHTTALSLPACPEESPLLVGPMLIEFNMPVDLELVAKQNPN
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complement(15755. .17099)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /evidence=not_experimental complement(14283) / gene="GGTA1"
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VGRCRMIRHSRDKKNEPNPQRFDRIAHTKETMLSDGLNSLTYQVLDVQRYPLYTQITV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DTIFNRAKLLNVGFQEALKDYDYTCFVFSDVDLIPMNDHNAYRCFSQPRHISVAMDKF
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24065. .24252,38835. .:
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complement(14305. .14310)
                                                                                                                                                                                                                                                                                    /note="AluJb repeat: matches 1.
24409. .24625
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19877. .20157
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                                                                                                                                                                                                                                                                                                                                                  /note="AluJo repeat:
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91.9%;
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                                                                                                                                                                                             Score 117.4;
Pred. No. 5.3
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                                                                                                                                                                                                                                                                                                        .305 of consensus"
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Grand-Pierre, N., Grant, G., Hagos, B., Heaford, A., Horton, L.,
Howland, J.C., Iliev, I., Johnson, R., Jones, C., Kann, L., Karatas, A.,
Klein, J., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J.,
Levine, R., Lieu, C., Liu, G., Locke, K., Macdonald, P., Marquis, N.,
McCarthy, M., McEwan, P., McGurk, A., McKernan, K., McPheeters, R.,
Meldrim, J., Meneus, L., Mihova, T., Miranda, C., Mienga, V., Morrow, J.,
Murphy, T., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P.,
O'Neil, D., Olivar, T. M., Oliver, J., Peterson, K., Pierre, N.,
Pisani, C., Pollara, V., Raymond, C., Riley, R., Rogov, P., Rothman, D.,
Roy, A., Santos, R., Schauer, S., Severy, P., Spencer, B.,
Stange-Thomann, N., Stojanovic, N., Subramanian, A., Talamas, J.,
Tesfaye, S., Theodore, J., Tirrell, A., Travers, M., Trigilio, J.,
Vassiliev, H., Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W.J.,
Voung, G., Zainoun, J., Zimmer, A. and Zody, M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (01-APR-2000) Whitehead Institute/MIT Center Research, 320 Charles Street, Cambridge, MA 02141, USA (bases 1 to 176584)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Birren,B., Linton,L., Nusbaum,C., Lander,E., Abraham,H., Allen,N., Anderson,S., Baldwin,J., Barna,N., Bastlen,V., Beda,F., Boukhiy,L., Boukhgalter,B., Brown,A., Burkett,G., Callymore,A., Castle,A., Choepel,Y., Colangelo,M., Collins,S., Collymore,A., Cooke,P., DeArellano,K., Dewar,K., Diaz,J.S., Collymore,A., Cooke,P., DeArellano,K., Dewar,K., Diaz,J.S., Collymore,A., Cooke,P., DeArellano,K., FitzHugh,M., Gage,D., Collymore,A., Doyle,M., Ferreira,P., FitzHugh,M., Gage,D., Collymore, Collymore, Collymore, Cooke,P., Doyle,M., Ferreira,P., FitzHugh,M., Gage,D., Collymore, Collymore, Collymore, Collymore, Cooke,P., Doyle,M., Ferreira,P., FitzHugh,M., Gage,D., Collymore, Collymor
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Homo sapiens chromosome 9 clone
SEQUENCE, 29 unordered pieces.
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HTG; HTGS_PHASE1; HTGS_DRAFT
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Sequencing vector: M13; M77815; 100% of reads
Sequencing vector: M13; M77815; 100% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 162821 bases at least Q40
Consensus quality: 162946 bases at least Q30
Consensus quality: 172552 bases at least Q20
Insert size: 182000; agarose-fp
Insert size: 173784; sum-of-contigs
Quality coverage: 4.0 in Q20 bases; agarose-fp
Quality coverage: 4.2 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NOTE: This is a 'working draft' sequence. It currently consists of 29 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
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g of 6286 bp in length
of 100 bp
g of 7596 bp in length
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of 8591 bp
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f of 4478 bp
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g of 3006 bp :
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of 2510 bp
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of 361 bp in length
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42878. .49203
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29361, .33317
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24783. 29260
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18050. .21350
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14944. .17949
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21451. .24682
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/chromosome="9"
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/mol_type="genomic DNA"
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38419. .42777
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gap of 100 bp
contig of 12196 bp i
gap of 100 bp
contig of 10556 bp i
gap of 100 bp
contig of 10556 bp i
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Pred. No. 5.3e-27;
0; Mismatches 11
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49303 57894 57994 64071 64171

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136201 66 0 14944 17950 18050 21351 21451 24683

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chemistry: Dye-primer ET; 100% of reads Chemistry: Dye-terminator Big Dye; 0% of reads Assembly program: Phrap; version 0.990319 Consensus quality: 179431 bases at least 040 Consensus quality: 180878 bases at least 030 Consensus quality: 181543 bases at least 020 Consensus quality: 181543 bases at least 020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
Submitted (04-APR-2000) Genome Sequencing Center, Washington Submitted (54-APR-2000) Genome Sequencing Center, Washington Submitted (54-APR-2000) Genome Sequencing Center, Washington St. Fig. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insert size: 18300; agarose-fp
Insert size: 182699; sum-of-contigs
Quality coverage: 6.27 in Q20 bases; agarose-fp
Quality coverage: 6.20 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University school of Médicine, 4444 Forest Park Parkway, 8 MO 63108, USA On Aug 26, 2000 this sequence version replaced gi:9795673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequencing vector: plasmid; 0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequencing vector: M13; 100%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Web site:http://genome.wustl.edu/gsc/index.shtml
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 (bases 1 to 183099)
Waterston, R.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence of Homo sapiens clone Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                   as soon as it is available and the accession number will be preserved.
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HTGS_PHASE1; HTGS_DRAFT.
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                                                                                                                                Location/Qualifiers
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93301. .132512
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132613. .183099
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24414. .54520
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91.9%;
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Perfect score:
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Maximum DB seq
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                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                       Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2552756 seqs, 1349719017 residues
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180
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1: /SIDS1/gcgdata/g.
2: /SIDS1/gcgdata/g
                                                                                                                                                                                                                                                                                                                                                                                               JSIDS1/gcgdata/geneseq/geneseqn-emb1/NA1989.DAT:*
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/SIDS1/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT:*
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  ABV86895
ABV89019
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AAT64673
AAZ77490
AAC98176
AAH34805
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                                                                                                                                                                                                   Description
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                  n colon cancer
n colon cancer
n tumour-assoc
n pancreatic c
n PEC-60-type
n ovarian tumo
n colon cancer
  colon
cancer
cancer
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colon canc	ABQ60575	24	362		6.	5
င်ဒ	AAN80460	9	341		σ.	44
BamHI	AAQ78611	15	302		ς.	43
hPSTI.M2 gene for	AAT84523	18	238		σ,	42
	Φ	12	234	0	6.	41
encoding	AAN91165	10	234		6.	40
Pancreas secretory	AAQ14261	12	200		36.6	39
Lambda phage cII s	. AAN91164	10	200		6.	38
Synthetic DNA enco	AAN82222	9	182		6.	37
creatio	AAN81535	9	179		6.	36
huma	AAQ01577	11	171	20.3	36.6	35
DNA	AAN82230	9	182	20.7	7.	34
DNA	AAN82229	9	182		37.2	ω
DNA	AAN82223	9	182	20.8	37.4	32
1c DNA	AAN82226	9	182	0	37.4	3
DNA	AAN82225	9	182	0	37.4	30
Synthetic DNA enco	AAN82224	9	182		37.4	29
Lung cancer	ABL65420	24	368	1.	8	28
Lung cancer rela	ABL64781	24	368		38.2	27
Gene #3850 usec	ABN97352	24	368		œ	26
Human pan	ABK86355	24	368	1.	8	25
Synthetic	AAN82221	ٯ	182	21.2	8	24
Modified	AAQ01494	Н	171	۲.	8	23
Synthetic DNA	AAN82233		182	1.		22
Synthetic DNA	AAN82228		182	1.	8	21
Synthetic DNA enc	AAN82227		182	1	8	20
pSKAN8 fragment #	AAT15982	17	415	۲.	9	19
Human pan	AAQ12109	2	314	22.1	9.	18
Synthetic DNA	AAN82232		182		9	17
	AAN82217	ø	182		9	16
n pla	AAQ24083	13	482	ω.		15
	AAN82216	9	182	u	41.4	14
Ω	AAN80030	9	182	ω	41.4	13
	AAN80452	9	168	ω	41.4	12
ce enco	AAN80453	9	168	4	44.6	1
colon	ABQ60577	24	283	76.3	137.4	10
Human colon cancer	ABQ60075	24	468	84.3	151.8	9

ALIGNMENTS

RESULT 1 ABV8695/c ID ABV86895 standard; cDNA; 323 BP.

ABV86895;

13-DEC-2002 (first entry)

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06-FEB-2001;
28-MAR-2001;
10-JUL-2001;
WPI; 2002-608400/65.
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                                                                                         Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human colon cancer related cDNA SEQ ID NO 206
                                                                                                                                                                                  (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-NOV-2001; 2001WO-US43704.
                                                                                         Xu J, Chenault RA,
                                                                                                                                                                                                                                                                           ; 20000$-25222P.
; 20010$-267011P.
; 20010$-279670P.
; 20010$-304837P.
                                                                                         Meagher MJ,
                                                                                         Secrist H,
                                                                                         King
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New isolated tumor diagnosis, preventi

prevention

and/or

polynucleotide and polypeptide, i/or treatment of cancer, in par

tide, useful particular

colon

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RESULT 2
ABV89019
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Best Local S
Matches 180
                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention are useful for the diagnosis, prevention and/or treatment of cancer, particularly colon cancer. (I) can be used in gene therapy and (I) and (II) are useful in pharmaceutical compositions such as vaccines. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (I) encoding a polypeptide (II, ABP67991-ABP67996) comprising: (i) any comprements of (i); (iii) at least 20 contiguous residues of (i); (iv) sequences that hybridize to (i), under moderately stringent conditions; (v) sequences having at least 75% or 90% identity to (i); or (vi) degenerate variants of (i). The compositions and methods of the present reputation and refer the formula of the present reputation.
     New isolated
                           WPI; 2002-608400/65.
                                                   Stolk
                                                                                            20-NOV-2000; 2000US-252222P.
06-FEB-2001; 2001US-267011P.
28-MAR-2001; 2001US-279670P.
10-JUL-2001; 2001US-304037P.
                                                                                                                                                                             01-AUG-2002.
                                                                                                                                                                                                    WO200258534 -
                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                      13-DEC-2002
                                                                                                                                                      19-NOV-2001;
                                                                                                                                                                                                                                                                                                                             ABV89019;
                                                                                                                                                                                                                                                                                                                                                  ABV89019 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                       (CORI-) CORIXA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention relates to a human colon tumour expressed polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                    121 CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGCCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  217
                                                JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             180;
                                                                                                                                                                                                                                                                                colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                            97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                colon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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                                                                                                                                                                                                                                                                                                                                                                                                         CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Хu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
   tumor colon polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 66 A; 81 C;
                                                                                                                                                      2001WO~US43704
                                                                                                                                                                                                                                                                                                     (first
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                                                                                                                                                                                                                                                         cancer; cytostatic;
                                                                        CORP
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                                                Chenault
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                                                                                                                                                                                                                                                                              related cDNA SEQ
                                                                                                                                                                                                                                                                                                                                                  cDNA;
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                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      266pp +
                                                                                                                                                                                                                                                                                                                                                  323
                                                RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93 G; 83 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 180; DB 24; Pred. No. 1.1e-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                               Meagher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence Listing;
                                                                                                                                                                                                                                                        tumour; gene therapy; vaccine;
                                                                                                                                                                                                                                                                                ID NO
   and
                                                                                                                                                                                                                                                                                2330
polypeptide, useful for
                                                Secrist
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                               King
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(I) encoding a polypeptide (II, ABP67991-ABP67996) comprising: (i) any of CC 2600 fully defined nucleotide sequences (ABP8669-ABP89289); (ii) CC complements of (i); (iii) at least 20 contiguous residues of (i); (iv) csquences that hybridize to (i), under moderately stringent conditions; CC (v) sequences having at least 75% or 90% identity to (i); or (vi) cd degenerate variants of (i). The compositions and methods of the present cancer, particularly colon cancer. (I) can be used in gene therapy and CC (I) and (II) are useful in pharmaceutical compositions such as vaccines. CC Note: The sequence data for this patent did not form part of the printed cospecification, but was obtained in electronic format directly from WIPO cat fip.wipo.int/pub/published_pct_sequences.
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AAV38073
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Best Local :
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                                                                                                                                                                                                  CDS
                                                                                                                                                                                                                                                       Human; tumour-associated Kazal inhibitor; TAKI; inflammation; cancer; ulcerative colitis; Crohn's disease; inflammatory cytokine; elastase; serine protease; glycoprotein hormone; trypsin; chymotrypsin; pepsin;
                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                14-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                              AAV38073 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis,
                                                                               06-NOV-1996;
                                                                                                      05-NOV-1997;
                                                                                                                              14-MAY-1998
                                                                                                                                                     WO9820132-A1
                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 323 BP; 83 A; 93 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                       (INCY-) INCYTE PHARM INC
                                                                                                                                                                                                                                                                                                                                                       WV38073;
                                                                                                                                                                                                                                     sapiens.
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                                                                                                                                                                                                                                                                                                       thmour-associated Kazal inhibitor encoding cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevention and/or treatment of cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                               96US-0744670
                                                                                                      97WO-US20204.
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                                                                                                                                                                                                 Location/Qualifiers 34..294
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                                                                                                                                                                           /product=
                                                                                                                                                                                                                                                                                                                                                                              cDNA;
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                                                                                                                                                                          "tumour-associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 180;
Pred. No.
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                                                                                                                                                                                                                                                        pepsin;
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WPI; 1998-286948/25 P-PSDB; AAW62074.

Bandman O,

Goli SK,

Murry

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ABV97573
ID ABV97
XX ABV ABV9
XX ABV ABV9
XX Huma
XX Huma
XX WO2(
XX OW)
AN OO-1
AN OO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 180
                                                                                                                                                                                                                        31-JAN-2001;
09-FEB-2001;
21-MAR-2001;
28-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TAKI). TAKI is expressed in response to inflammatory cytokines and functions in the inhibition of serine proteases or glycoprotein hormones present in diseased tissue. The TAKI can be used to inhibit the tissue destruction associated with the production of excess proteases such as trypsin, chymotrypsin, elastase, or pepsin. Products from the present invention can be used in the treatment of e.g. proliferative cell division in inflamed intestinal tissues, ulcerative colitis, Crohn's disease or cancer. The products can also be used for detection and
                                              Benson
                                                                                                                                                                                                                                                                                                              30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                            30-JAN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                        08-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                     W0200260317-P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; pan
cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human par
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 -JAN 2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  АВУ9₹573;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV97573 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5; Fig 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated tumour-associated Kazal inhibitor, TAKI - used develop products for the diagnosis and treatment of e.g. inflammation, ulcerative colitis, Crohn's disease or cancer
                                                                                      (CORI-)
                                                                                                                                                                              16-MAY-2001;
12-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens
  2002-627435/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    180;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ц
                                                                                      CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ncreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          389 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      creas; car;
tumour;
                                            Kalos MD,
                                                                                                                               2001US-265305P
2001US-265682P
2001US-278631P
2001US-278631P
2001US-278132P
2001US-291633P
2001US-305484P
2001US-313999P
2001US-333626P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                            2002W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Spp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer; gene therapy; vaccine;
ur; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          105 A; 105 C; 102 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer expressed cDNA SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA; 394
                                                                                                                                                                                                                                                                                                                                                          -US02781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          encodes human tumour-associated Kazal inhibitor
                                            Lodes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 180; DB\
Pred. No. 1.2e-
; Mismatches
                                         Rersing DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9;
                                         Hepler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunostimulant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                            WI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            389;
                                            Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       171
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RESULT 5
AAT64673
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cc any of a group of over 4000 nucleotide sequences (ABV94628-ABV99145);

(c) by complements of (a); (c) sequences consisting of at least 20

cc contiguous residues of (a); (d) sequences that hybridize to (a), under

cc contiguous residues of (a); (d) sequences that hybridize to (a), under

cc contiguous residues of (a); (d) sequences that hybridize to (a), under

cc contiguous residues of (a); (e) sequences having at least 73% or 90%

cc defect cancer in a patient and compositions comprising polypeptides,

cc polynucleotides, antibodies, fusion proteins, T cell populations and

antigen presenting cells expressing the polypeptide are useful in

cc treating pancreatic cancer and stimulating an immune response. The

cc polynucleotides can be used as probes or primers for nucleic acid

cc hybridisation, in the design and preparation of ribozyme molecules for

cc inhibiting expression of the tumour polypeptides and proteins in the

cumour cells, in vaccines and for gene therapy.

cc specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 180
                                                                                                                                CDS
                                                                                         sig_peptide
                                                                                                                                                                                                                                                                                                                                AAT64673 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID
                          WO9715596-A1
                                                               mat_peptide
                                                                                                                                                                      Homo
                                                                                                                                                                                               antigen;
                                                                                                                                                                                                           Human; PEC-60-type protein; digestive disorder; immune disorder;
nervous disorder; gastrointestinal tract hormone; gastric tissue
                                                                                                                                                                                                                                                  Human PEC-60-type protein encoding
                                                                                                                                                                                                                                                                               22-DEC-1997
                                                                                                                                                                                                                                                                                                       AAT64673;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pancreatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotide and pancreatic for diagnosing, preventing and/or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to
                                                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                                    197
                                                                                                                                                                                                                                                                                                                                                                                                                           121
                                                                                                                                                                                                                                                                                                                                                                                                                                                       137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             180;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                              CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCXTGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAA&ACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                               antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
                                                                                                                                                                                                                                                                            (first entry)
                                                                                        /*tag= a
/product=
43..120
                                                               /*tag= k
121..300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO 2981; 300pp + Sequence Listing; English.
                                                                                                                              Location/Qualifiers 43..303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 115
                                                 /*tag=
                                                                                                                   /*tag=
                                                                                                                                                                                               gastrointestinal tract hormone; gastric
                                                                                                                                                                                                                                                                                                                                cDNA to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 102 C;
                                                                             ٥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         an
                                                                                                     PEC-60-type_protein
                                                                                                                                                                                                                                                                                                                                mRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         isolated polynucleotide (I) comprising:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 180; DB 24;
Pred. No. 1.2e-51;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100 G;
                                                                                                                                                                                                                                                                                                                                  398
                                                                                                                                                                                                                                                   CDNA.
                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumor polypeptides,
cancer, particularly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       part of the printed directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        394;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (a)
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                                                                                                                                                                                                                                                                                                                                                                                                    256
                                                                                                                                                                                                                                                                                                                                                                                                                              180
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01-MAY-1997

22-OCT-1996;

96WO-JP03061.

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RESULT 6
AAZ77490
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                                                                                                                                                                                                                                                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence encodes a human PEC-60-type protein. The protein is a gastrointestinal tract hormone secreted by the gastric tissues. It is useful for treatment or diagnosis of digestive, immune or nervous disorders, or as an antigen to raise antibodies against them. Large amounts of the protein may be produced by using the corresponding cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PEC-60-type protein - immune or nervous disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-258958/23. P-PSDB; AAW15067.
                             WPI; 1999-591920/51
                                                                                   09-APR-1998;
                                                                                                      09-APR-1998;
                                                                                                                         21-OCT-1999.
                                                                                                                                            DE19817557-A1.
                                                                                                                                                             Homo sapiens
                                                                                                                                                                                          Expressed sequence
                                                                                                                                                                                                            Human
                                                                                                                                                                                                                               10-APR-2000
                                                                                                                                                                                                                                                                  AAZ77490 standard;
 New nucleic
                                               Rosenthal A,
                                                                                                                                                                                                                                                 AAZ77490;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 398 BP; 107 A; 110 C; 105 G; 76 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kamata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SAGA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                               180;
                                                                                                                                                                               therapy;
                                                                                                                                                                                                                                                                                                                  241
                                                                                                                                                                                                                                                                                                                                    121
                                                                                                                                                                                                                                                                                                                                                       181
                                                                                                                                                                                                                                                                                                                                                                                           121
                                                                                                                                                                                                            ovarian
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                  AAY76598,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SAGAM
                                                                  METAGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                    GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT 180
                                                                                                                                                                                                                                                                                                                                                                                                           GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 10;
acid
                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                               Specht T,
                                                                                                                                                                               treatment;
                                                                                                                                                                                                                             \first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CHEM
                                                                                                      98DE-1017557
                                                                                    98DE-1017557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95JP-0280272.
                                                                 GES GENOMFORSCHUNG MBH.
                                                                                                                                                                                                         mor cDNA library derived EST
sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17pp; Japanese.
                   AAY76599.
                                                                                                                                                                                         tag; EST; human; ovarian tumor; anticancer;
                                                                                                                                                                                                                                                                  cDNA; 402
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sekine-S,
                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                SS
                                               Hinzmann
expressed
                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for
                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 180; DB 18;
Pred. No. 1.2e-51;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yamaguchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment
in
                                               Schmitt
ovarian,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Т;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or diagnosis of digestive,
                                                                                                                                                                                                          fragment 41.
                                              Ą
and
                                              Pilarsky
                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
some
other,
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                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                               Dahl
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cancer
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円 (7)

(HUMA-) HUMAN GENOME SCI INC

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RESULT 7
AAC98176
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Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            밁
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence tags) from a particular tissue type before comparison of the expression patterns. This allows a significantly longer fragment of the gene to be revealed, so should reduce the number of failures associated with the fact that ESTs from different libraries may represent different parts of the same unknown gene, distorting the estimated frequency of occurrence in a particular tissue. AAZT7450-Z77572 represent the human ovarian tumor cDNA library derived EST fragments described in the method of the invention and encode the protein fragments represented in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    [This invention describes novel nucleic acid (cDNA) sequences (A) which have anticancer activity and are highly expressed in ovarian tumor tissue (and some also in testis and breast cancer tissue). The products of the invention can be used for gene therapy. (A) are used (i) for recombinant expression of polypeptides (B) and (ii) to isolate complete genes. (B) are used (i) to identify agents suitable for treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                         09-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ovarian cancer; (ii) directly for treating this form of cancer (including expression from gene therapy vectors) and (iii) for generation of specific antibodies. (A) are identified by assembling ESTs (expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tissues, and derived polypeptides, for treatment of ovarian cancer and identification of therapeutic agents \,
                                                                                       08-MAR-2000;
                                                                                                                                                                                                                                                                       neural disorder; immune system disorder; muscular disorder
                                                                                                                                                                                                                                                                                            nephrotropic;
                                                                                                                                                                                                                                                                                                                          Human; colon cancer; colon cancer antigen; diagnosis; detection; identification; cytostatic; cardioactive; neuroprotective; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC98176 standard; cDNA; 433
                                                   12-MAR-1999;
                                                                                                                           21-SEP-2000
                                                                                                                                                               WO200055351-A1
                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                     reproductive disorder; gastrointestinal disorder; renal disorder; infectious disease; cardiovascular disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                   Human colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 402 \pp; 108 A; 111 C; 107 G; 76 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY76505-Y76638.
                                                                                                                                                                                                                                                                                                               immunomodulatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                   cancer antigen nucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                         2000WO-US05883
                                                                                                                                                                                                                                                                                          ory; muscular; gynaecological; gastrointestinal;
antiinfective; antibacterial; gene theraby; wound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       170;
                                                     99US-0124270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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Pred. No. 1.2e-51;
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                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID
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243

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RESULT 8
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KW CO1(
XXX Hom
KW CO1

XXX HOm
XXX WO2
PD 05-
XXX PF 28-
XXX 29-
PR 29-
PR 03-
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Best Local S
Matches 180
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                                                                                                                                                                                                       05-APR-2001
                                                                                                                                                                                                                                                      WO200122920-A2
                                                                                                                                                                                                                                                                                                        Homo
                                                                                                                                                                                                                                                                                                                                                          colorectal cal
                                                                                                                                                                                                                                                                                                                                                                                  Human; colon
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human co.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-SEP
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                                                                         29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective, immunomodulatory, gynaecological, gastrointestinal, vulnerary, nephrotropic, antiinfective and antibacterial activities,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC97991 to AAC98063 encode the human colon cancer associated proteins, called human colon cancer antigens, given in AAB53234 to AAB54006. The human colon cancer antigens can have cytostatic, cardioactive, muscular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Colon cancer associated gene sequences, referred to antigens, useful for the treatment, prevention, and
                          (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                   28-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                               cancer antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                   2000WO-US2652
                                                                                                                                                                                                                                                                                                                                                                                  cancer; colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10; 2104pp; English
                                                                         99US-0157137.
99US-0163280.
                                                                                                                                                                                                                                                                                                                                                          inoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA; 433 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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                                                                                                                                                                                                                                                                                                                                                          SS
                                                                                                                                                                                                                                                                                                                                                                               cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                            encoding cDNA SEQ ID NO:1887
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 21;
2e-51;
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RESULT 9
ABQ60075
ABQ60075
AC ABQ6
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps. Additionally, N may be used to produce the colon cancer-associated Ps. By inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and N can be used in the prevention, diagnosis and treatment of colorectal carchomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                       02-OCT-2000; 2000US-237271P
                                                                                                        02-OCT-2001;
                                                                                                                                                               11-APR-2002
                                                                                                                                                                                                                                                                                                                    genetic
                                                                                                                                                                                                                                                                                                                                                  Human;
                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABQ60075
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N.B. Pages 666 to 682 and page 7053 of the missing at time of publication, meaning no SEQ ID NO:1027 to 1052, 7921 and 7922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer-associated nucleic acid molecules (N) the proteins are collectively known as colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH32943 to AAH37195
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                                                                                                                                                                                                              WO200229086-A2
                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABQ60075;
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                                                                                                                                                                                                                                                                                                                                                                                                 colon cancer related nucleotide sequence SEQ ID NO:3770
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                                                                                                                                                                                                                                                                                                                    colon cank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      preven
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                                                                                                        2001WO-US30732
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                       diagnostic; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA; 468 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                  cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Birse CE,
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                                                                                                                                                                                                                                                                                                                                                tissue profiling; forensic; mapping;
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Pred. No. 1.\
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ξe-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequences are present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and proteins (P), where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CACATGGTAGAGTCTCCAACCTGT
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180

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(FARB) BAYER CORP

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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cc expressed in cancer tissues. ABB78993 to ABB79004 represent proteins cc encoded by the ABQ60787 to ABQ60787 nucleic acid sequences. (I) can be cused in antisense therapy. An antibody immunoreactive with a polypeptide cc encoded by (I) is useful for detecting cancer in a patient sample, and cc for detecting the presence of absence of a polynucleotide encoded by a cc nucleic acid which hybridises to (I) in a cell. A probe/primer derived cf from (I) can be used for determining the presence of a nucleic acid which hybridises to (I) in a cell. A probe/primer derived comparison to used for determining the presence of cc list from a patient. (I) is useful for determining the presence of cc colon cancer in a cell or tissue type, for determining the presence or cc state of other type of cancer, in antisense therapy, to generate cc corresponding gene resides, and in tissue profiling, forensics, genetic canalysis, mapping and diagnostic applications. (I) can be used to raise antibodies, and to screen for peptide analogues and antagonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Sim
Matches 175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleuc acid that is differentially expressed tissues useful for determining the presence of colon cancer or tissue type, and in antisense therapy
               02-OCT-2001; 2001WO-US30732
                                             11-APR-2002
                                                                                                                                    genetic
                                                                                                                                                   Human;
                                                                                                                                                                                                                                           ABQ60577;
                                                                                                                                                                                                                                                                        ABQ60577 standard; cDNA; 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Fig 1; 796pp;
                                                                        WO200229086-A2
                                                                                                                                                                            Human colon cancer related nucleotide sequence SEQ ID NO:4272.
                                                                                                                                                                                                             02-AUG-2002
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                                                                                                                                                  colon cancer;
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                                                                                                                                    analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 96.775; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCCAGATGTCCAACCTGGTCTGCGGCACTGATGGGCTCACATATAC-GAATGAATGCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTCTGCTTGGCCCGGATAAAAACCAACNAGGACATCCAAGATCATGAAAGATGGCAAATG
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                                                                                                                                                                                                            (first entry)
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                                                                                                                                    diagnostic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84.3%;
96.7%;
                                                                                                                                                  cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 151.8; DB Pred. No. 6e-42; O; Mismatches
                                                                                                                               tissue profiling; forensic; mapping; antisense therapy; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Catino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , LT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABQ56306 to ABQ60787 represent isolated nucleic acids (I) differentially expressed in cancer tissues. ABB78993 to ABB79004 represent proteins encoded by the ABQ60776 to ABQ60787 nucleic acid sequences. (I) can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1;
                               05-MAY-1988
                                                                                         Homo sapiens
                                                                                                                        Human pancreatic secretory trypsin inhibitor; pancreatic proteolytic zymogens.
                                                                                                                                                                      Sequence encoding novel human pancreatic secretory trypsin inhibitor (HPSTI).
                                                                                                                                                                                                                      25-MAR-2003
17-SEP-1990
                                                                                                                                                                                                                                                                      AAN80453;
                                                                                                                                                                                                                                                                                                   AAN80453 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or tissue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-426115/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCCANATGTCCAACCTGGTNTGCGGCACTGATNGGCTCACATATACAAATGAATGCCAG
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                                                                                                                                                                                                                                                                                                                                                                                   CTCTGNTTGG-CCGNATAAAAACCAAACAGNACATCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     283 BP; 67 A; 84 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fig 1; 796pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         type, and in antisense therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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(first entry)
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                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.3%;
94.3%;
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                                                                                                                                                                                                                                                                                                   168
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09-OCT-1987;

87WO-US02585

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Result
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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Match
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-021-298-55
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US-09-312-283C-357
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US-09-149-933-2
US-09-149-933-362
US-09-149-933-362
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ULT 2 08-744-670-2 atent No. 5858710 GENERAL INFORMATION APPLICANT: Bandm APPLICANT: MUTTY APPLICANT: MUTTY TITLE OF INVENTION NUMBER OF SEQUENCI CORRESPONDENCE ADD ADDRESSEE: Inc:	1 1 61 61 121	I 1 -065-019-	20 20 20 20 20 20 20 20 20 20 20 20 20 2
JT 2 JURNES 2, Application US, Lent No. 5858710 EENT NO. 5858710 SINERAL INFORMATION: APPLICANT: Bandman, Ol, APPLICANT: Goli, Surya APPLICANT: Murry, Lynn TITLE OF INVENTION: TUI NUMBER OF SEQUENCES: 6 CORRESSONDENCE ADDRESS: ADDRESSEE: Incyte Pha	GGAAAGCTC GGAAAGCTC TCCCAGATG TCCCAGATG CTCTGCTTG	Separation sap	15.7 15.7 15.7 15.7 15.7 14.8 14.8 14.8
ion US/087, ii Olga an, Olga Surya K. , Lynn E. , Lynn E. TUMOR- ES: 6 ES: 6 DRESS: yte Pharma	CCTTTCTCI	US/09 i Tomoke ngo yu Man PE T-US MBER: 998-04 3 3 2.0 100.0	466 4 771 2 771 4 2495 4 2495 2 2525 2 2526 3 2226 4 2226 4 11529 3 11529 4 11529 4 11
/08744670 ga K. E. MOR-ASSOCIATED	GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT	LIGN IKE 065,	US-09-617-1 US-08-972-1 US-09-267-1 US-09-117-1 US-09-617-1 US-08-972-1 US-08-972-1 US-08-972-1 US-08-8112-1 US-08-8112-1 US-09-252-1
KAZAL	NTCTGTGAAC.	PROTEIN AND 019	-804-5 2-008-3 -027-1 -804-1 -008-1 -260A-1 -626A-1 -626A-1 -754A-3 -748-1 -991A-9494 -991A-9494 -312D-893
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US-09-149-933-2
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                                                                                                                                                                                                                                                     Sequence 2, Application Patent No. 5958699
GENERAL INFORMATION:
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             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COmpatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                             APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
APPLICANT: Murry, Lynn E.
TITLE OF INVENTION: TUMOR
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LIBRARY: SINTTUT01
CLONE: 1539065
                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Diskett
                                                                                                                                                                               NUMBER OF SEQUENCES:
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                                                                                                               STREET: 3174 Por
CITY: Palo Alto
                                                                                       COUNTRY:
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STATE: CA
                                                                                                        STATE:
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                                                                                                                                    E: Incyte Pharmaceuticals, 3174 Porter Drive
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FastSEQ Version 2.0
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US-09-065-019-2
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 388 base pairs
LENGTH: 380 bace pairs
TYPE: nucleic acid
                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 398
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Patent No. 6280968
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                            Matches
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CURRENT FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 3
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: HUMAN PEC-60-LIKE PROTEIN AND DNA ENCODING THIS PROTEIN FILE REFERENCE: 6700PCT-US
                                                                                                                                                                                                                                                                                                                APPLICANT: Sekine, Shingo APPLICANT: Kamata, Kouju
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
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ATTORNEY/AGENT INFORMATION:
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  181
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1 GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT 60
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36,749
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Pred. No. 4.5e-56;
Pred. No. 4.5e-56;
                                                                                                         Score 180; DB 3;
Pred. No. 4.5e-56;
Mismatches 0;
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; ORGANELLE:
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US-08-880-829-21
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US-08-880-829-21
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                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: (212) 426767
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Roettgen, Peter
TITLE OF INVENTION: A Collection of Phagemids, A
TITLE OF INVENTION: Collection of Escherichia Coli
TITLE OF INVENTION: Cells Carrying The Phagemids, A
TITLE OF INVENTION: Collection of Phagemid Particles
From OF INVENTION: Produced From Said Collection
Produced From Particles
                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 687-6000
TELEPHAX: (212) 682-3485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM-XT COMPATIBLE
OPERATING SYSTEM: DOS 3.3:
SOFTWARE: WORDDERFECT 5.0
CURRENT APPLICATION DATA:
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                            MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 0
FILING DATE: 06/02/95
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MEDIUM TYPE: 3-1/2" DISKETTE
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                                                                                                                                                                                                                                                              ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Joseph T. Eisele
ADDRESSEE: Kane, Dalsimer, Sullivan,
ADDRESSEE: Levy, Eisele and Richard
STREET: 711 Third Avenue
CITY: New York
STATE: New York
COUNTRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/880,829 FILING DATE: 23-JUN-1997 CLASSIFICATION: 435
                                                                                                                                        INDIVIDUAL ISOLATE:
DEVELOPMENTAL STAGE:
                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: EISELE, JOSEPH T
REGISTRATION NUMBER: 25
                                                              CELL LINE:
                                                                              CELL TYPE:
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                                                                                                   TISSUE TYPE:
                                                                                                                     HAPLOTYPE:
                                                                                                                                                                                                ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 458 base pairs
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10017-4059
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5925559
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US-08-235-515A-26
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                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                            FEATURE
                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 302 base pairs
                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
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                                                                                              LOCATION: 1..302
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                NAME: Murphy Jr., Gerald M. REGISTRATION NUMBER: 28,977 REFERENCE/DOCKET NUMBER: 11
                                                                                                                                                                                                                                                            TELEPHONE: /UJ - 100:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Birch, Stewart, Kolasch & Birch STREET: P.O. Box 747 CITY: Falls Church
                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/235,515A FILING DATE: 29-APR-1994
LOCATION:
              NAME/KEY:
                                         LOCATION:
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                                                                                     OTHER
                                                                                                                                         NAME/KEY:
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5840518
                                                                                   INFORMATION:
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27..89
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57.7%;
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                                                                                Figure
                                                                                  /label= polynucleotide
/note= "j-xl-y-z(psti),
Figure 5"
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Pred. No. 6.2e-05;
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US-08-235-515A-26
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Patent No. 650093
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Best Local Similarity
      Matches
                                 Query Match
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APPLICANT: Janice
APPLICANT: Jeffre
                                                                                                                                                                                                TELEFAX: (650) 845-4166 INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/016,434
FILING DATE: HEREWITH
                                                                                                                                                                                                                          REGISTRATION NUMBER: 37,071
REFERENCE/DOCKET NUMBER: 97
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 855-0555
                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION NUMBER: 37,
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Jeffrey J. Seilhamer
TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF SIGNALING
TITLE OF INVENTION: PATHWAY GENE EXPRESSION
                                                                                                        IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                   Local
                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Jeffrey J. Seilhamer
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Score 36.6; DB 4;
Pred. No. 0.00089;
0; Mismatches 54;
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LENGTH: 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/221,298
CURRENT FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 112
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Xu, Jiangchun TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER FILE REFERENCE: 210121.471
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PEATURE:
NAME/KEY: modified_base
                           LOCATION: (385)
OTHER INFORMATION: Where
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NAME/KEY: modified_base
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OTHER INFORMATION: Where
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OTHER INFORMATION: Where
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OTHER INFORMATION: Where
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OTHER INFORMATION: Where n
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                     FEATURE:
                                                     NAME/KEY: modified_base
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NAME/KEY: modified_base
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Title: Perfect score: Sequence:

Scoring table:

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Minimum DB Maximum DB

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length: length:

Sequence 68711, A
Sequence 11934, A
Sequence 12055, A
Sequence 19567, A
Sequence 53, Appl
Sequence 51, Appl
Sequence 151, App
Sequence 111, App
Sequence 19737, A
Sequence 19737, A
Sequence 19749, A
Sequence 11448, A
Sequence 16448, A
Sequence 16448, A
Sequence 1648, A
Sequence 6164, Appl
Sequence 6619, App
Sequence 6619, App
Sequence 7086, App
Sequence 7086, App
Sequence 7086, App
Sequence 19370, A

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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US-60-485-150-1215
US-60-493-007-3005
US-60-500-315-1122
US-60-500-315-11231
US-60-500-315-11538
US-09-947-914-75
US-60-487-610-243
PCT-US02-36071A-139
US-60-487-610-1955
US-60-487-610-1955
US-60-487-610-1955
US-60-487-610-196487
US-60-485-450-315-116487
US-60-485-450-371
US-60-485-450-371
US-60-485-450-171
US-60-487-610-1923
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11539, App
1243, App
13550, A
19550, A
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US-60-487-610-19749/c

Sequence 19749, Application US/60487610

GENERAL INFORMATION:

APPLICANT: CARGILL, Michele
APPLICANT: HUNNI, Hongjin

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CL001469

CURRENT APPLICATION NUMBER: US/60/487,610

CURRENT FILING DATE: 2003-07-17

NUMBER OF SEQ ID NOS: 97101

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 19749

LENGTH: 28753

TYPE: DATE: 28753
                                                                                                                                            RESULT 2
US-60-485-450-12159/c
• Sequence 12159, Application US/60485450
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Best Local Sim:
Matches 124;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C
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                  APPLICANT: CARGILL, Michele
APPLICANT: CHANG, Sheng-Yung
APPLICANT: CHANG, Sheng-Yung
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: RESPONSE TO INTERFERON TREATMENT IN H
TITLE OF INVENTION: THRECTED SUBJECTS, METHODS OF D
TITLE OF INVENTION: THRECOF
FILE REFERENCE: CL001470
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7 US-60-500-315-12055
7 US-60-500-315-12055
7 US-60-500-337-19567
6 US-09-947-914-53
6 US-09-947-914-151
7 US-60-487-610-19737
7 US-60-487-610-19737
7 US-60-487-610-19749
7 US-60-485-450-12159
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5 US-10-653-047-6619
5 US-10-425-114A-7086
7 US-60-500-337-19370
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Pred. No. 2.5e-31;
0; Mismatches 11
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                                                   ; OTHER INFORMATION: Genbank Accession No. M64780
US-60-493-007-3005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-60-493-007-3005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA; Homo sapiens US-60-485-450-12159
                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION
Best Local Similarity
                                                                                                                                                             SEQ ID NO 300
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/493,007
CURRENT FILING DATE: 2003-08-07
PRIOR APPLICATION NUMBER: US 60/353,171
PRIOR FILING DATE: 2002-02-04
PRIOR APPLICATION NUMBER: US 60/363,534
PRIOR FILING DATE: 2002-03-13
                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/394,253 PRIOR FILING DATE: 2002-07-09
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BQ ID NO 12159
LENGTH: 28753
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                                                                                                                                                                              SOFTWARE:
                                                                                                                                                                                            Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 3518
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Primary rat hepatocyte toxicity modeling FILE REFERENCE: 44921-5113-01-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Mendrick, Donna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                        TYPE: DNA
                                                                                       ORGANISM: Rattus norvegicus FEATURE:
                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                   FILING DATE: 2002-04-19
APPLICATION NUMBER: US 60/374,139
FILING DATE: 2002-04-22
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                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 60/371,150 FILING DATE: 2002-04-10
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/373,601
                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 60/371,413 FILING DATE: 2002-04-11
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/370,248 FILING DATE: 2002-04-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/371,135
FILING DATE: 2002-04-10
APPLICATION NUMBER: US 60/371,134
                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 2002-04-10
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Elashoff, Michael
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91.9%;
17.0%;
73.6%;
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Pred. No. 2.5e-31;
0; Mismatches 11;
Score 30.6; D
Pred. No. 1.6;
                                                                                                                                                                                                           See File Wrapper or
                DB 7;
              Length 7286;
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APPLICANT: CARGILL, Michele
TITLE OF INVENTION: POLYMORPHISMS IN NUCLEIC
TITLE OF INVENTION: ENCODING HUMAN G-PROTEI
TITLE OF INVENTION: OF DETECTION AND USES T
FILE REFERENCE: CL001484
CURRENT APPLICATION NUMBER: US/60/500,315
CURRENT FILLING DATE: 2003-09-05
NUMBER OF SEO ID NOS: 69978
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11231
LENGTH: 203938
TYPE: DNA
ORGANISM: Homo sapiens
US-60-500-315-11231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: POLYMORPHISMS IN NUCLEIC TITLE OF INVENTION: ENCODING HUMAN G-PROTEI TITLE OF INVENTION: OF DETECTION AND USES THE REFERENCE: CLOO1484

CURRENT APPLICATION NUMBER: US/60/500,315
CURRENT FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 69978
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SEQ ID NO 11222
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TYPE: DNA
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US-60-500-315-11231
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                                                                    176355
                                                                                                                                         176295
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                                                                                   89 CTGATGGGCTCACATATACGAATGAATGCCAGCTCTGCTTGGCCCGGATAAAAACCCAAAC 148
                                                                                                                                                             29 TCTGTGAACACATGGTAGAGTCTCCAACCTGTTCCCAGATGTCCAACCTGGTCTGCGGCA 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 GAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT 60
                                                                  CAGCTTTCTGTACAGTGTTATTTGCAAGCCCACTAAGCTGGACAGGGAAAAAGAACCTGAC
                                                                                                                                       TCTGGCTACCCATGGCAGCCCAGGAAAGAGAGTCCTGGATGCACAGGAGCTTCTCTGGGT 176354
                               AGGACATCCAGATCATGAA
                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            POLYMORPHISMS IN NUCLEIC ACID MOLECULES
ENCODING HUMAN G-PROTEIN COUPLED RECEPTOR PROTEINS, METHODS
OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              POLYMORPHISMS IN NUCLEIC ACID MOLECULES
ENCODING HUMAN G-PROTEIN COUPLED RECEPTOR PROTEINS, METHODS
OF DETECTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Michele
                                                                                                                                                                                                                              16.8%;
51.1%;
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69.5%;
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RESULT 6 US-60-500-315-11538/c

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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

SOURCE ORGANISM ORGANISM REFERENCE AUTHORS TITLE JOURNAL	LOCUS DEFINITION ACCESSION VERSION KEYWORDS	RESULT 1 AR166138		44	c 42 43	44	. 3 G	37	ယ ယ တ ဟ	33 4	32	30	28 29	. c 26	N	c 23	22	, N	c 19	70	15			0 د			ວ າ ປ	ı.e. u	221	Result No.
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	æ	g the same	encodin	Human PEC-60-like protein and DNA encoding the same	50-like prot	Human PEC-6	TITLE
		ta, K.	nd Kama	Kato, S., Yamaguchi, T., Sekine, S. and Kamata, K.	amaguchi,T.,	Kato, S., Ya	AUTHORS
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                                E 1 (bases 1 to 358)

Kato,M., Yamaguchi,T., Sekine,S. and Kamata,M.

KHUMAN PEC-60LIKE PROTEIN AND DNA CAPABLE OF COI

Patent: JP 1997124698-A 2 13-MAY-1997;

SAGAMI CHEM RES CENTER

OS HOMO Sapiens (human)

PN JP 1997124698-A/2

PD 13-MAY-1997

PF 27-OCT-1995 JP 1995280272

PI KATO MASASHI, YAMAGUCHI TOMOKO, SEKINE SHI

MITSUHISA

CO7K14/575,C07H21/04,C12N15/09//A61K38/22,

PC (12P21/02,C12R1:19);

CC strandedness: Double;

CC topology: Linear;

FH Key

FT Source

/organism='Homo sapiens'

FT Source

/tissue_type='stomach canc
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                                                                                                                                 Human nucleic acid sequences from ovarian tumour tissue patent: WO 9953040-A 41 21-CCT-1999;
SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (DE), BERND (DE); ROSENTHAL ANDRE (DE); METAGEN GES FUER GENO (DE); FILARSKY CHRISTIAN (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
/111 c 107 g 7
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                                                                                                      Location/Qualifiers
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             Score 398;
Pred. No.
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No. 2.8e-99;
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Indels

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Gaps

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RESULT 4
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TITLE
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Best Local Similarity
Matches 388; Conser
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Sequence
AR028526
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Bandman,O., Goli,S.K. and Murry,L.E.
Tumor-associated kazal inhibitor-like
                                                                                                                                                                                                                                                                               polynucleotides
Patent: US 5858
                                                                                                                                                                                                                                                                                                                                         Unknown
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                             CTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGTTCCCAGATGT
                                                                                                                     TGGCCTTGGCCGCCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCC
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                                                                    CTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGTTCCCAGATGT
                                                                                                           TGGCCTTGGCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCC
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ilarity 100.0%;
Conservative
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Location/Qualifiers
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105 c 102 g
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5 5858710.
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1.6e-96;
hes 0;
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                                      RESULT 6
BD007521
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AUTHORS
TITLE
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FEATURES
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AR075817
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Tumor-associated KAZAL t
BD007521.1 GI:18635894
JP 2001503629-A/1.
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Sequence 2 :
AR075817
AR075817.1
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I (bases 1 to 388)

Bandman,O., Goli,S.K. and Murry,L.E.

Tumor-associated Kazal inhibitor

Patent: US 5958699-A 2 28-SEP-1999;

Location/Qualifiers
  unidentified
unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                             ACATGAAATAAAAGATCCAGCCCAACTG 398
                                                                                                                                                            GGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGGAGAGGATATG
                                                                                                                                                                                                CCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGCTGATCCCACA
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Pred. No.
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240 250

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RESULT 7
AF048700
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Homo sapiens
AF048700
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PN JI
PD 21
PF 05
PR 06
PR 06
PC CC
PC C1
PC C12N15
PC A6
FT SC
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Bandman,O., Goli,S.K. and Murry,L.E.
Tumor-associated KAZAL type inhibitor
Patent: JP 2001503629-A 1 21-MAR-2001;
INCYTE PHARMACEUTICALS INC
                          Homo sapiens
              Eukaryota;
                                       Homo
                                                              AF048700.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                            GAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGGAGAGAGTATG
                                                                                                                                                                                                                         GGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGTGGGCATGGAGAGATATG
                                                                                                                                                                                                                                                           CTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGTTCCCAGATGT 190
                                                                                                                                                                                                                                                                                                                                                                                                                        TGGCCTTGGCTGCCCTTGTTGTGGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCC 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGGGTAATCGCCC
                                                                                                                                                                                      ACATGAAATAAAAGATCCAGCCCAACTG
                                                                                                                                                                                                                                                                                  CCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGCTGATCCCACA 310
                                                                                                                                                                                                                                                                                                          CTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCCAACCTGTTCCCAGATGT
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                                     sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAR-2001
05-NOV-1997 JP 1998521770
06-NOV-1996 US 08/744670
01-NOV-1996 US 08/744670
01-A BANDMAN, SURYA K GOLI, LYNN E MURRY
C12N15/09, A61K38/55, A61K39/395, A61P1/00, A61P35/00, C07K14/81,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
source
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.5/00,C12N5/00,
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Location/Qualifiers
1...388
 ; Metazoa;
Eutheria;
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
105 c 102 g 7
                                                             GI:2935439
                                                                                    386 l
gastrointestinal
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 Chordata;
Primates;
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1. .388
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Pred. No. 1.
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l peptide (PEC-60) mRNA,
Craniata; Ve Catarrhini;
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1.6e-96;
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Vertebrata; Euteleostomi;
i; Hominidae; Homo.
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RESULT
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AUTHORS
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KEYWORDS
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Sus scrofa (pig)
Sus scrofa
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Eutheria; Cetartiodactyla; Suina; Suidae;
1 (bases 1 to 364)
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Direct Submission
Submitted (17-FEB-1998) Shanghai Institute
Submitted (18-FEB-1998) Shanghai 200031, China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human gastrointestinal Unpublished 2 (bases 1 to 386)
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PEC-60=gastrointestinal
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Location/Qualifiers
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llarity 100.0%;
Conservative
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/protein_id="AAC05124.1"
/db_xref="GI:2935440"
/translation="MANCOWIALALALLYVDREVPVAAGKLPFSRMPICEHMVESP
TCSQMSNLVCGTDGLTYTNECQLCLARIKTKQDIQIMKDGKC"

97 c 98 g 72 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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12. .272
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| peptide
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                                        Euteleostomi;
Sus.
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                                    pec-60 gene.
Sus scrofa (pig)
Sus scrofa
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Molecular cloning of PEC-60 and expression of its mRNA and per the gastrointestinal tract and immune system J. Biol. Chem. 267 (28), 19829-19832 (1992)

93015834
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entry [NCBI gibbsq 115614] from the original journal article.
This sequence comes from Fig. 1.
Location/Qualifiers
Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 364)
                                                                         x67109.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                    GGCACTGATGGGCTCACATATACGAATGAATGCCAGCTCTGCTTGGCCCGGGATAAAAAACC
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                                                                                                                                                                                                                                                                                                                                                                                            CTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCCCTTTCTCAAGAATG 144
                                                                                                                                                                       AAACAGGACATCCAGATCATGAAAGATGGCAAATGCTGATCCCACAGGAGCACCTCAAGC
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23, 202
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                                                                         GI:2033
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                                                                                 mpgc60 gene.
Mus musculus (house mouse)
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
 Unpublished 2 (bases 1
                    Krause,R., Hemberger,M., Messerschmid,M., Kothary,R. an Molecular cDNA cloning and expression of murine Mpgc60, predominantly expressed in the intestinal tract
                                                                                                                                                                      MMMPGC60
                                                                     Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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to 515)

Euteleostomi;
; Murinae; Mus

and Fundele, R

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AUTHORS
TITLE
               M.musculus mRNA for Mpgc60 Y11505
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Submitted (22-JUN-1992) M. Metsis, Dept. of Medical Chemistry, Lab. of Molecular Neurobiology, Karolinska Institute, Box 604 00, 10401
Y11505.1 GI:2344863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Metsis, M.
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/translation="MAVRLWVVALALAALFIVDREVPVSAEKQVFSRMPICEHMTESP
DCSRIYDPVCGTDGVTYESECKLCLARIENKQDIQIVKDGEC"
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93 c 108 g 72
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/product="peptide PEC-60"
/protein_id="CAA47482.1"
/db_xref="GI:2034"
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/db_xref="taxon:9823"
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Pred. No. 6.2e-58;
0; Mismatches 62
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Best Local Sim:
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Direct Su
1 (bases 1 to 180)
Kato,S., Yamaguchi,T., Sek
Human PEC-60-like protein
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Sequence 1 from patent US
AR166137
                                                           Unknown
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                                           Unclassified
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Ihnestr. 73, 14195 Berlin,
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/protein_id="CAA72282.1"
/db_xref="GI:2344864"
/db_xref="SPTREMBL:035679"
/db_xref="SPTREMBL:035679"
/translation="MANHIWLVTLTLGFLLGMDRELMVSAGSLVFPRMPFCEHMAELP
NCPQTPNLICGTDGLTYENECHLCLTRMKTMKDIQIMKDGQC"
129 c 136 g 95 t
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/db_xref="taxon:10090"
/tissue_type="intestine"
/dev_stage="adult"
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/mol_type="mRNA"
/strain="NMRI"
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75.3%;
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 Sekine, S. and Kamata,
in and DNA encoding t
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6280968.
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121 GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTGTCTCCAACCTGT 180
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PD JP 19
PD 13-MA
PF 27-OC
PF KATO
OMITSUHISA
PC C07K1
PC C12P2
PC C12P2
PC C12P2
PC C5TPA
CC topol
FH Key
FH Sourc
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HUMAN PEC-60LIKE PROTEIN AND DNA CAPABLE OF
Patent: JP 199124698-A 1 13-MAY-1997;
SAGAMI CHEM RES CENTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 180)
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E13090.1 GI:32519
JP 1997124698-A/1.
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13-MAY-1997
27-OCT-1995 JP 1995280272
77-MTO MASASHI, YAMAGUCHI TOMOKO, SEKINE SHINGO,
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JP 1997124698-A/1
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                                       Conservative
                                                                                                                                                                                                                                source
                                                                                                                                                                                                                                                                     topology: Linear;
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Location/Qualifiers
1. .180
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                                                                                                    /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
49 c 40 g 3
                                                                                                                                                               /product='PEC-60-like protein'
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49 c 40 g
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100.0%; Pred. No. 7.5e-39;
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Pred. No. 7.5e-39;
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                                                                                                                                                                                                       /tissue_type='stomach
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PEC-60-like
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GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT

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Maximum DB seq
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Perfect score:
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                    349.4
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length: 2000000000
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99.2
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398
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Listing first 45 summaries
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/SIDS1/gcgdata/geneseq/geneseqn-embl/NA1995.DAT:*
/SIDS1/gcgdata/geneseq/geneseqn-embl/NA1995.DAT:*
/SIDS1/gcgdata/geneseq/geneseqn-embl/NA1996.DAT:*
/SIDS1/gcgdata/geneseq/geneseqn-embl/NA1999.DAT:*
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/SIDS1/gcgdata/geneseg/geneseqn-emb1/NA2001B.DAT:*
/SIDS1/gcgdata/geneseg/geneseqn-emb1/NA2002.DAT:*
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                  AAT64673
AAZ77490
AAC98176
AAH34805
AAW38073
ABV97573
ABQ60075
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                                                                          colon cancer
                                                                                                                 PEC-60-type
ovarian tumo
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pancreatic c
colon cancer
                    colon cancer
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Human prostate		23	904	٠	40.6	
Human prostate exp		23	692	10.2	40.6	
Human prostate		23	692	0	40.6	
Human		20	608	0		
Human pancreatic		21	604	0		
Human		25	432	0		
Human		24	379	0		
Human		24	362	0	40.6	
Human pancreatic s		9	341	0	40.6	
	AAN82223	9	182	10.2	40.6	
Synthetic DNA		9	182	0	40.6	
		9	182	0	40.6	
Synthetic		9	182	0	40.6	
		11	171	0	41.2	
		13	482	0	41.4	
		17	415	0	41.4	
S		9	182	10.4	41.4	
		9	168	10.4	41.4	
Synthetic DNA		9	182	0	42	
		9	182	0	42	
Synthetic DNA		9	182	0	42	
Lung cancer		24	368	0		
Lung cancer rela		24	368	10.6	42.2	
Gene #3850 used		24	368	0	42.2	
Human pancreas se		24	368	0	Ñ	
Human pancreatic s		12	314	0	42.4	
Synthetic DNA		9	182	0		
Synthetic DNA		9	182	0	43	
	AAN82216	9	182			
Synthetic	AAN80030	9	182	1	44.6	
Sequence encodin	AAN80453	Ģ	168	-	4	
Human secreted	AAC04139	21	621	-	45.6	
Human secreted	AAC04140	21	573	Ϊ.	Ġ	
Human	ABN39626	24	60	5		
Human colon	565	24	166	Ģ	01.	
Human colon cance	ABQ60577	24	283	64.0	254.8	
Human colon cancer	90	24	323	0	21.	

ALIGNMENTS

RESULT 1

AAT64673;

AAT64673 standard; cDNA to mRNA; 398 BP

22-DEC-1997 (first entry)

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AAT64673
ID AAT6
XX AAT6
AC AAT6
XX LAT6
XX LA
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                                                      22-OCT-1996;
                                                                                                                                                   01-MAY-1997.
                                                                                                                                                                                                                                                                                                                                                                                mat_peptide
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121..300
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RRESULT 2
AAZ7740
ID AAZ77
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AC. AAZ7
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DT 10-2
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EXP
EXP
KW Exp
KW Exp
KW gene
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GS Home
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PN DE19
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence encodes a human PEC-60-type protein. The protein is a gastrointestinal tract hormone secreted by the gastric tissues. It is useful for treatment or diagnosis of digestive, immune or nervous disorders, or as an antigen to raise antibodies against them. Large amounts of the protein may be produced by using the corresponding cDNA.
                                            Homo
                                                                                                                                                 AAZ77490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PEC-60-type immune or nervous
 21-OCT-1999
                       DE19817557-A1
                                                                  gene
                                                                              Expressed
                                                                                                                          10-APR-2000
                                                                                                                                                                     AAZ77490 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SAGA ) SAGAMI CHEM
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DB; AAW15067.
                                           sapiens
                                                                 therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTAATCGCCCTGGCCTTGCCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCA
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                                                                                                                                                                                                                                                                                                                                                                            GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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                                                                                                                                                                                                                                                                                              TGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGG
                                                                                                                                                                                                                                                                                                                    CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                GCAGGCCCCAGCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
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                                                                                                                                                                                                                                                                        TGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGG
                                                                              sequence tag; EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kato S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
                                                                                                                         (first
                                                                  treatment;
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disorders
                                                                                                                         entry)
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                                                                                                  cDNA library derived EST
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Pred. No. 1.2
0; Mismatches
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1.2e-115;
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                                                                              tumor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC This invention describes novel nucleic acid (cDNA) sequences (A) which CC have anticancer activity and are highly expressed in ovarian tumor CC tissue (and some also in testis and breast cancer tissue). The products CC of the invention can be used for gene therapy. (A) are used (i) for CC recombinant expression of polypeptides (B) and (ii) to isolate complete CC genes. (B) are used (i) to identify agents suitable for treatment of CC ovarian cancer; (ii) directly for treatment suitable for cancer (CC (including expression from gene therapy vectors) and (iii) for generation CC of specific antibodies. (A) are identified by assembling ESTs (expressed CC sequence tags) from a particular tissue type before comparison of CC expression patterns. This allows a significantly longer fragment of the CC gene to be revealed, so should reduce the number of failures associated CC with the fact that ESTs from different libraries may represent different CC parts of the same unknown gene, distorting the estimated frequency of CC courrence in a particular tissue. AAZ77450-Z77572 represent the method CC varian tumor CDNA library derived EST fragments described in the method CC of the invention and encode the protein fragments represented in
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY76505-Y76638.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tissues,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              identification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid sequences expressed in ovarian, and some other, cosues, and derived polypeptides, for treatment of ovarian cancer
                           361
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364
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                                                                                                                                                                                                                                                                GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTAACACATGGTAGAGTCTCCCAACCTGT 180
                                                                                                                                                                                                                                                                                                                              GTAATCGCCCTGGCCTTGGCTGCCCTTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCA 120
                                                                                                                                                                                                                                                                                                                                                                                            GCAGGCCCCAGCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
                                                        TGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGG 360
                                                                                                                      GTAATCGCCCTGGCCTTGCCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCA
                                                                                                                                                                                                                                                                                                                                                                          GCAGGCCCCAGCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    402 BP;
           AGAGGATATGACATGAAATAAAAGATCCAGCCCAACTG
                                                                                                                                                     CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                   GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Pred. No. 1.2e-115;
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RESULT

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Best Local :
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                                                                                                                                                                                                                                                                             neuroprotective, immunomodulatory, gynaecological, gastrointestinal, vulnerary, nephrotropic, antiinfective and antibacterial activities, and can be used in gene therapy. The colon cancer antigen polynucleotides, proteins and antibodies to the proteins are useful for the prevention, treatment and diagnosis of colon disorders, such as colon cancer. The polynucleotides may be used in diagnostics and research, such as for chromosome identification, and as hybridisation probes. The proteins may also be used to prevent diseases such as neural disorders, immune system disorders, muscular disorders, reproductive disorders, gastrointestinal disorders, wounds, renal disorders, infectious diseases, and cardiovascular disorders. AAC98764 to AAC98772 and AAB54077 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC97991 to AAC98763 encode the human colon cancer associated proteins, called human colon cancer antigens, given in AAB53234 to AAB54006. The human colon cancer antigens can have cytostatic, cardioactive, muscular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Colon cancer associated gene sequences, referred to as colon cancer antigens, useful for the treatment, prevention, and diagnosis of colon disorders such as colon cancer -
                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 610; 2104pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neural disorder; immune system disorder; muscular disorder;
reproductive disorder; gastrointestinal disorder; renal disorder;
infectious disease; cardiovascular disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nephrotropic; antiinfective;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-MAR-2001
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DB; AAB53419.
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                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                        433
GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCCAACCTGT
                                                                GTAATCGCCCTGGCCTTGGCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCA
                                                                                                           GCAGGCCCCAGCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
                                                                                                                                          GCAGGCCCCAGCCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
                                               GTAATCGCCCTGGCCTTGCCCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCA
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                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                      B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                muscular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cdna;
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                                                                                                                                                                                                                                    A; 113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gynaecological; gastrointestinal;
; antibacterial; gene therapy; wou
tem disorder; muscular disorder;
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                                                                                                                                                                                                                                      C; 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cardioactive; neuroprotective; vulnerary;
                                                                                                                                                                                     Score 395;
Pred. No.
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                                                                                                                        AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon cancer associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps. by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carrinoms and cancers. AAH37196 to AAH37204
present invention. N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present for
                                                                                                   and AAB77789 represent sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding 4277 human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human colon cancer antigen encoding cDNA SEQ ID NO:1887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 3403; 9803pp; English
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99US-0163280.
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and/or treating colorectal cancers -
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                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                       Human; tumour-associated Kazal inhibitor; TAKI; inflammation; cancer; ulcerative colitis; Crohn's disease; inflammatory cytokine; elastase; serine protease; glycoprotein hormone; trypsin; chymotrypsin; pepsin;
                                                                                                                                                                                                                                                                                                  Human tumour-associated Kazal inhibitor encoding cDNA
          P-PSDB;
                                           Bandman O,
                                                                                       06-NOV-1996;
                                                                                                             05-NOV-1997;
                                                                                                                                   14-MAY-1998
                                                                                                                                                                                                                                                                                                                             14-SEP-1998
                                                                                                                                                                                                                                                                                                                                                   AAV38073;
                                                                                                                                                                                                                                                                                                                                                                       AAV38073 standard;
                                                                 (INCY-)
                                                                                                                                                        WO9820132-A1
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hes 395;
                    1998-286948/25
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          AAW62074
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                                                                                                                                                                                                                                                                                                                                                                                                                                       AGAGGATATGACATGAAATAAAAGATCCAGCCCAA 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTCTGCTTGGCCCGGATAAAAACCCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAAGATGGCAAATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCAGGCCCCAGCCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGG
                                           Goli SK,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP;
                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                 PHARM
                                                                                       96US-0744670
                                                                                                            97WO-US20204
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/product=
                                                                                                                                                                                                   Location/Qualifiers 34..294
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100.0%;
                                           Murry LE;
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                                                                                                                                                                              "tumour-associated Kazal
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                        ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated tumour-associated Kazal inhibitor, TAKI - used develop products for the diagnosis and treatment of e.g. inflammation, ulcerative colitis, Crohn's disease or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 389
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362
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                            ACATGAAATAAAAGATCCAGCCCAACTG
                                                                                     GGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCCATGGAGAGGATATG
                                                                                                                    CCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGCTGATCCCACA
                                                                                                                                                                                                                                                                                                      TGGCCTTGGCTGCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCC 130
                                                                                                                                                                                                                                                                                                                                                                  GCCAGCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGGGTAATCGCCC
ACATGAAATAAAAGATCCAGCCCAACTG
                                                         GGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGGAGAGGATATG
                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        105 A; 105 C; 102 G; 76 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            97.5%; Score 388; DB 19; 100.0%; Pred. No. 1.7e-112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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XX ABY9752
XX ABY99
XX I4-J
XX Huma
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XX EYTC
XX EYTC
XX HOMC
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XX WO20
XX WO3-J
XX WO3-J
XX WO3-J
                                                                                                                  14-JAN-2003
                                                                                                                                   ABV97573;
                                                                                                                                                  ABV97573 standard;
                                                                                                                 (first entry)
                                                                                                                                                  cDNA; 394
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Human pancreatic cancer expressed CDNA SEQ ID NO 2981

tuman; pancreas;
cytostatic; tumou tumour; cancer; gene therapy;
ur; gene; ss. vaccine; immunostimulant;

sapiens

WO200260317-A2

08-AUG-2002.

30-JAN-2002; 2002WO-US02781

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC The invention relates to an isolated polynucleotide (1) comprising: (a) CC any of a group of over 4000 nucleotide sequences (ABV94628-ABV99145); CC (b) complements of (a); (c) sequences consisting of at least 20 cc contiguous residues of (a); (d) sequences that hybridize to (a), under CC moderately stringent conditions; (e) sequences having at least 75% or 90% cidentity to (a); or (f) degenerate variants of (a). Polypeptides (c) identity to (a); or (f) degenerate variants of (a). Polypeptides cc (ABP6859-ABP68637) encoded by (I) and oligonucleotide can be used to (c) detect cancer in a patient and compositions comprising polypeptides, polynucleotides, antibodies, fusion proteins, T cell populations and cc treating pencreatic cancer and stimulating an immune response. The cc treating pencreatic cancer and stimulating an immune response. The cc polynucleotides can be used as probes or primers for nucleic acid composition, in the design and preparation of ribozyme molecules for c inhibiting expression of the tumour polypeptides and proteins in the cc tumour cells, in vaccines and for gene therapy.

CC Note: The sequence data for this patent did not form part of the printed content with our content and enterence of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the printed content of the pri
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Matches 350
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31-JAN-2001; 2001US-265682P.
09-FEB-2001; 2001US-267568P.
21-MAR-2001; 2001US-278651P.
28-APR-2001; 2001US-287112P.
16-MAY-2001; 2001US-291631P.
12-JUL-2001; 2001US-39394P.
20-AUG-2001; 2001US-3333626P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide and pancreatic tumor polypeptides, for diagnosing, preventing and/or treating cancer, particularly pancreatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 394 BP; 115 A; 102 C; 100 G; 73 T; 4 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1;
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                                                                                                                                                                     AGAGTCTCCAACCTGTTCCCAGATGTCCAACCTGGTCTGCGGCACTGATGGGCTCACATA
                                                                                                                                                                                                                                                                                                                                                            AGTGCCAGTGGCAGCAGGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGT 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO
ACAGTGGTGGGCATGGAGAGGATATGACATGAAATAAAAGATCCAGCCCAA
                                                                                GAAAGATGGCAAATGCTGATCCCACAGGAGCACCTCAAGCCATGAAGTGTCAGCTGGAGA
                                                                                                                                                                                                               TACGAATGAATGCCAGCTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCAT
                                                                                                                                                                                                                                                                                                                                           AGTGCCAGTGGCAGCAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGT
                                                                                                                                                                                                                                                                                                                                                                                                                          GGCCGTCCGCCAGTGGATAATCGCCCTGGCCTTGGCTGCCCTTGTTGTTGTGGACAGGGA
                                                                                                                                                                                                                                                      AGAGTCTCCAACCTGTTCCCAGATGTCCAACCTGGTCTGCGGCACTGATGGGCTCACATA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87.8%;
99.7%;
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Pred. No. 2.8e-100;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Burgess C, Astle JH, Thiaglingam A, Lewis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibodies, and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 analysis; diagnostic; antisense therapy;
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                      TCCCAGATGTCCAACCTGGTCTGCGGCACTGATGGGCTCACATATAC-GAATGAATGCCA
                                                                                                                                                                                                                                                                                                                                                        GCAGGCCCCAGCCAGCCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
                                                                                                                                                        GGAAAGCTCCCTTTCTCAAGAATGCCCCATCTGTGAACACATGGTAGAGTCTCCAACCTGT
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                                                                                                        NGAAAGCTCCCTTTCTCAAGAATGCCCATCTGTGAACACATGGTATAGTCTCCAACCTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative.
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No. 8.7
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RESULT 8
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Query Match
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                                                       (I) encoding a polypeptide (II, ABP67991-ABP67996) comprising: (i) any c 2600 fully defined nucleotide sequences (ABV8669-ABV89289); (ii) complements of (i); (iii) at least 20 contiguous residues of (i); (iv) sequences that hybridize to (i), under moderately stringent conditions; (v) sequences having at least 75% or 90% identity to (i); or (vi) degenerate variants of (i). The compositions and methods of the present invention are useful for the diagnosis, prevention and/or treatment of cancer, particularly colon cancer. (I) can be used in gene therapy and (II) and (II) are useful in pharmaceutical compositions such as vaccines. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at figure 1.
                                                                                                                                                                                                                                                                                                                                                                                        20-NOV-2000;
06-FEB-2001;
28-MAR-2001;
                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                      diagnosis,
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                                                                                                                                                                                                                                                                       prevention
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; 2001US-267011P.
; 2001US-279670P.
; 2001US-304037P.
                                              BP;
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d/or treatment of ca
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ancer, in particular
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GCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGGGTAATCGCCCTGGC

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RESULT 9
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The invention relates to a human colon tumour expressed polynucleotide (I) encoding a polypeptide (II, ABP67991-ABP67996) comprising: (i) any c 2600 fully defined nucleotide sequences (ABV8669-ABV89289); (ii) complements of (i); (iii) at least 20 contiguous residues of (i); (iv) sequences that hybridize to (i), under moderately stringent conditions: (v) sequences having at least 75% or 90% identity to (i); or (vi) degenerate variants of (i). The compositions and methods of the present invention are useful for the diagnosis, prevention and/or treatment of cancer, particularly colon cancer. (I) can be used in gene therapy and (II) and (II) are useful in pharmaceutical compositions such as vaccines. Note: The sequence data for this patent did not form part of the printed
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06-FEB-2001;
28-MAR-2001;
                                                                                                                                                                                                                                                                                                                                               diagnosis,
cancer -
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                                         Unpublished Contact: John Quackenbush Contact: John Quackenbush The Institute for Genomic Research 9712 Medical Center Dr., Rockville, Tel: 301 838 3528 Fax: 301 838 0208
                                                                                                       Quackenbush, J.

Assessment of gene expression patterns in a model of metastasis using a 19,200 element cDNA microarray
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                                                                                                                                                                     Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                      Email: johng@tigr.org
Plate: 149
  Seq primer: Reverse.
Location/Qualifiers
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                                                                 Email: yongsung@mail.kribb.re.kr
Plate: 51 row: E column: 10
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l: +82-42-860-4470
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Pred. No. 1.1e-101;
; Mismatches 0;
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                                                                                                                       of Bioscience & Biotechnology
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J.W., Bouffard,G., Smith,D. and Peterson,K. Expressed sequence tag analysis of human retina for Project: Retbindin, an abundant, novel retinal cDNA splicing of other retina-preferred gene transcripts Mol. Vis. 8 (4), (2002) In press
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BQ636767
BQ636767.1 GI:21761226
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Wistow,G., Bernstein,S.L., Wyatt,M.K.,
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/clone="$14K402-51-E10"
/cell_line="K402"
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Pred. No. 1.3e-89;
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 graeme@helix.nih.gov
01 row: b column:
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="he01b10"
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/dev_stage="Adult"
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                                                                      321 bp mRNA linear EST 21-AI NCI_CGAP_CO3 Homo sapiens cDNA clone IMAGE:926191 SW:PE60_PIG P37109 PEPTIDE PEC-60 PRECURSOR. ;, ml
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Insert Length: 417 Std Error: 0.00
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Contact: Robert Strausberg,
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Arraying: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Preparation: M. Bento Soares, Ph.D
                                                                                                CTCAAGCCATGAAGTGTCAGCTGGAGAACAGTGGTGGGCATGGAGAGGATATGACATGAA 377
                                                                                                                                                                                                                                                                                                                                                                                           GCTGCCCTCCTTGTTGTGGACAGGGAAGTGCCAGTGGCAGCAGGAAAGCTCCCTTTCTCA 138
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/Clone_lib="NCI_CGAP_CO3"
/note="vector: pT7T3D-Pac (Pharmacia) with a modified
/note="vector: pT7T3D-Pac (Pharmacia) with a modified
/note="vector: Site_1: Rot I; Site_2: Ecc RI; lst strand cDNI
/note="vector: Site_1: Ecc RI; lst strand cDNI
was prepared from 12 pooled bulk tumor samples and primed
with a Not I - oligo(dT) primer. Double-stranded cDNA was
ligated to Eco RI adaptors (Pharmacia), digested with Not
I and cloned into the Not I and Eco RI sites of the
modified pT7T3 vector. Library went through one round of
normalization. "
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/mol_type="mRNA"
/db_xref="taxon:9606"
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/lab_host="DH10B"
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/sex="pooled"
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wd. ET from
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NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
plate: LLAM8445 row: h column: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, | Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
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601070792F1 NIH_MGC_12
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                                                                                                                                                                                            GTAGAGTCTCCAACCTGTTCCCAGATGTCCAACCTGGTCTGCGGCACTGATGGGCTCACA 222
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                                           Athersys RAGE Library I
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/note="Organ: cervix; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT
Average insert size 1.4 kb. Library prepared by Life
Technologies."
300 c 247 g 88 t
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/clone="IMAGE:3456839"
/tlssue_type="cervical carcinoma cell line"
/lab_host="DH10B"
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/mol_type="mRNA"
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99.3%;
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Pred. No. 3.5e-69;
0; Mismatches 1
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282; Conservative
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1 (bases 1 to 537)

Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachadran, R., Whittington, J., Leventhal, C., Thornton, M., Boozer, S., Mays, R., Smith, E., Veloso, N., Kilka, A., Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danzig, J. and Ducar, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Creation of genome-wide protein expression libraries using random activation of gene expression nat. Biotechnol. 19 (5), 440-445 (2001)
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                 AA296922
EST112478 Retina
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Fax: 216 361 9596
Email: scain@athersys.com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact:
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/note="See 'Creation of Genome wide Protein Expression
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."
a 133 c 148 g 106 t
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No.
232 bp mRNA linear EST sapiens cDNA 5' end similar to PEC-60, mRNA sequence.
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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Maximum Match 100%
Listing first 45 summaries
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LENGTH: 398
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Best Local :
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NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin Ver. 2.0
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Sequence 2370, Ap	Sequence 2208, Ap	Sequence 2156, Ap	Sequence 362, App	Sequence 362, App	Sequence 17, Appl	Sequence 9, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 14, Appl	Sequence 6, Appli	Sequence 5, Appli	Sequence 4, Appli	Sequence 894, App	Sequence 357, App	Sequence 10, Appl

ALIGNMENTS

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Sequence 2, Application US/09065019 Patent No. 6280968 GENERAL INFORMATION:
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APPLICANT: Yamaguchi, Tomoko
APPLICANT: Sekine, Shingo
APPLICANT: Sekine, Shingo
APPLICANT: Kamata, Kouju
TITLE OF INVENTION: HUWAN PEC-60-LIKE PROTEIN
FILE REFERENCE: 6700PCT-US
CURRENT APPLICATION NUMBER: US/09/065,019
                                                                                                                                                                                                                                                                                                                                                                                                                               1 GCAGGCCCCAGCCTCAGGCTACACTATCCCAGGATCAGCATGGCCGTCCGCCAGTGG
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LIBRARY: SINTTI
CLONE: 1539065
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
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CITY: Palo Alto
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ADDRESSEE: Incyte Pharmaceuticals,
STREET: 3174 Porter Drive
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APPLICANT: Goli, Surya K.
APPLICANT: Murry, Lynn E.
APPLICANT: MURRY, Lynn E.
TITLE OF INVENTION: TUMOR-ASSOCIATED KAZAL INHIBITOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/7. FILING DATE: Filed Herewith PRIOR APPLICATION DATA:
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MEDIUM TYPE: Diskett
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NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
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                                                                                                                                                                                                                                                                                                                    Local
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Pred. No. 8.6e-118;
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INFORMATION FOR SEQ ID NO:
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NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,
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APPLICATION NUMBER: US/09/1-
FILING DATE: Filed Herewith
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LIBRARY: SINTTUT01
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LENGTH: 388 base pairs
TYPE: nucleic acid
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TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Diskett
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APPLICANT: MUXTY, LYND E.
TITLE OF INVENTION: TUMOR-ASSOCIATED KAZAL INHIBITOR
NUMBER OF SEQUENCES: 6
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LIBRARY: 51...
1539065
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                               CTTTCTCAAGAATGCCCATCTGTGAACACGTGTAGAGTCTCCCAACCTGTTCCCCAGATGT 180
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100.0%; Pred. No. 8.6e-118;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1
LENGTH: 180
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence 1, Application US/09065019 Patent No. 6280968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kato, Seishi
APPLICANT: Yamaguchi, Tomoko
APPLICANT: Sekine, Shingo
APPLICANT: Sekine, Shingo
APPLICANT: Kamata, Kouju
TITLE OF INVENTION: HUMAN PEC-60-LIKE PROTEIN AND DNA ENCODING THIS PROTEIN
FILE REFERENCE: 6700PCT-US
CURRENT APPLICATION NUMBER: US/09/065,019
CURRENT FILING DATE: 1998-04-17
NUMBER OF SEQ ID NOS: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                         Vatent No. 5925559
GENERAL INFORMATION:
                                                                                              APPLICANT: Collins, John
APPLICANT: Roettgen, Peter
TITLE OF INVENTION: A Collection of Phagemids, A
TITLE OF INVENTION: Collection of Escherichia Coli
TITLE OF INVENTION: Cells Carrying The Phagemids, A
TITLE OF INVENTION: Collection of Phagemid Particles
TITLE OF INVENTION: Produced From Said Collection
TITLE OF INVENTION: And Phagemid Particles
                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Joseph T. Eisele
                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                 ADDRESSEE:
                  ADDRESSEE:
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5925559
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                                                                                                                                                                                                                                                                                                                                                                                                                       CTCTGCTTGGCCCGGATAAAAACCAAACAGGACATCCAGATCATGAAAGATGGCAAATGC
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                Kane, Dalsimer, Sullivan, Kurucz,
Levy, Eisele and Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45.2%; Score 180; DB 3; 100.0%; Pred. No. 1.1e-49;
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                                            Sequence 1250, App
Patent No. 6500938
                                                                                                                                                                                                                                                                                                                                    Matches
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GENERAL INFORMATION:
APPLICANT: Janice
APPLICANT: Jeffre
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APPLICATION NUMBER: 08/458
FILING DATE: 06/02/95
APPLICATION NUMBER: Germar
FILING DATE: 06/07/94
ATTORNEY/AGENT INFORMATION:
NAME: EISELE, JOSEPH T.
REGISTRATION NUMBER: 25,33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212) 682-3485
TELEX: (212) 426767
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 10017-4059
COMPUTER READABLE FORM:
MEDIUM TYPE: 3-1/2" DISKETTE
COMPUTER: IBM-XT COMPATIBLE
OPERATING SYSTEM: DOS 3.3:
SOFTWARE: WORDPERFECT 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE: ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212) 687-6000
                                                                                                                                                                                                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
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nes 75; Conserv
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANELLE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INDIVIDUAL ISOLATE:
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                                                                                                                                                                                                                                                                        New York
Janice Au-Young
Jeffrey J. Seilhamer
                                                             Application US/09016434
                                                                                                                                                                                                                                                                                                                                     Conservative
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57.3%;
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                                                                                                                                                                                                                                                                                                                                     Score 41.4; DB 2;
Pred. No. 0.00056;
0; Mismatches 56;
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US-09-016-434-1250
                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-235-515A-26
                                                                                                                                                                                                                                                                                                                                                     Sequence 26, Application US/08235515A Patent No. 5840518
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APPLICANT: Morish
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Zeller, Karen J.
REGISTRATION UNMBER: 37,071
REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                              NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                             APPLICANT: NO. 5840: TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                   TITLE OF INVENTION:
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APPLICATION NUMBER:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS
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                                          ADDRESSEE:
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P.O.
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Kanamori, Toshinori
Kanamori, Toshinori
No. 5840518uhara, Masahiro
No. 5840518uhara, Masahiro
No. 5840518uhara, TRANSFORMANT TRANSFORMED WITH
TRANSFORMANT TRANSFORMENT, TRANSFORMANT TRANSFORMENT, TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT TRANSFORMANT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : (650) 855-0555
(650) 845-4166
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                                    Birch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INCYTE PHARMACEUTICALS, INC
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      Box 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HEREWITH
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                                Stewart,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1490
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                                    Kolasch & Birch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 432;
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                                                                                                                                VECTOR
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US-09-221-298-56/c
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                                                                                                                                                                                     RESULT 8
GENERAL INFORMATION:
APPLICANT: XU, Jiangchun
APPLICANT: XU, Jiangchun
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER
FILE REFERENCE: 210121.471
CURRENT APPLICATION UNMEER: US/09/221,298
CURRENT FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-235-515A-26
                                                                                                                                    Sequence 56, Application US/09221298 Patent No. 6284241
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 29-APR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Murphy Jr., Gerald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
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72; Conserv
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Virginia
                                                                                                                                                                                                                                   TGCTGA 296
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27..293
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90..293
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27..89
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/note= "j-xl-y-z(psti),
Figure 5"
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Pred. No. 0.0018;
Mismatches 5
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